

Result No.	Score	Query		Length	DB	ID	Description
		Match					
1	2687	51.7	4105	11	US-11-136-527-2482		Sequence 2482, A
2	416.8	8.0	600	11	US-11-136-527-6578		Sequence 6578, A
3	165.2	3.2	4053	11	US-11-091-883-433		Sequence 433, App
4	96.2	1.9	874	7	US-10-750-185-31648		Sequence 31648, A
5	96.2	1.9	874	7	US-10-750-623-31648		Sequence 31648, A
6	76.8	1.5	153376	11	US-11-121-086-5		Sequence 5, Appli
7	74	1.4	2170	7	US-10-750-185-31653		Sequence 31653, A
8	74	1.4	2170	7	US-10-750-623-31653		Sequence 31653, A
9	64.6	1.2	173602	11	US-11-121-086-25		Sequence 25, Appl
10	64.2	1.2	1400	11	US-11-136-527-7472		Sequence 7472, Ap
11	64.2	1.2	2380	11	US-11-136-527-3376		Sequence 3376, Ap
12	63.8	1.2	3456	11	US-11-136-527-155		Sequence 155, App
13	63.8	1.2	28536	11	US-11-011-3324-156		Sequence 156, App
14	63.2	1.2	4257	11	US-11-000-688-163		Sequence 163, App
15	62.8	1.2	2774	11	US-11-000-688-1278		Sequence 1278, Ap
16	59.8	1.2	2479	11	US-11-136-527-3373		Sequence 3303, Ap
17	59.8	1.2	241805	7	US-10-995-561-13215		Sequence 13215, A
18	58.4	1.1	415117	7	US-10-995-561-13274		Sequence 13274, A
19	58.2	1.1	3508	11	US-11-091-883-15		Sequence 15, Appl
20	58.2	1.1	3508	11	US-11-091-883-161		Sequence 161, App
21	58.2	1.1	26667	7	US-10-995-561-13375		Sequence 13375, A

241 ATATGGGGCGGTGGCTGCATGAATAGGACGGAGGATGGATCAGGTGGAGTTTTTCATA 300

QY	448	GCTATGAATCTTATCAAACTGAAGCTTAAAGGATATCAGTACCTCTCTGACCTTCCCTCT	507
Db	301	GCATGAAGCTCATCAAACTGAAGCTTGAAGGATATCAGTCCCTCCGCACTTCCCTCT	360
QY	508	GTCATGAAGACCAACAGTGTCTATTTCTAGCGCACAGCATTTCTGATGGAGGTATC	567
Db	361	GTCATGAAGACGAGCCAGCGGCACTCTCTAGTGCACAGCGTCTTGGTATAGAGGGATG	420
QY	568	GCCAGCATGCCACCGCTTACAGCTGTGTCTCAGTGCCCAATGGGATCCATTCAGTGTGT	627
Db	421	GCTGGAAATGCCACCACTGACAGCTGTGTCTCCGTGCCAATGGGCTCCATCCAGTGTGT	480
QY	628	GGAAATGTCCTCAACCTCTAGTATCTTCTGTCTCCACAGCAGCTGTGCCCCCTCTGGCTAAC	687
Db	481	GGAAATGTCCTCCGCTTAGTATCTTCTGTCTCCATCAGCAGCAGTGTCTCCCTGGCTAAC	540
QY	688	GGGGCTCCCCCTGTTATCAACCTCTGCTGCAATTTGCTCATCTCCAGCCATTTGCCA	747
Db	541	GGGGCCCCCTCTGTCTATACAGCTCTGCTGTGCAATTTGCTCATCTCCAGCCATTTGCCA	600
QY	748	AAGAGTCTTCTCTTTAGTAGATCTGGTCCAGGCTCAAACTAAACACTAAATTTACAAAAG	807
Db	601	AAGAGTCTTCTCTTACAGCATCTGGTCCAGGCTCAAACTAAACACTAAATTTACAAAAG	660
QY	808	GCACAGTCAATTTGATGTGGCAGTGTCCCAACAGTGGCAGAGTGGGCTGTCTCAGTCA	867
Db	661	GCACATCAATTTGATGTAGCAGCGCTCTGCAAGCGGCAAGTGGGCTGTGCTCAGTCTG	720
QY	868	TCAGAGCTGAATACAGGCAATTTATTTCAATAGTATGATGCAAACTATGATGAGCACTTA	927
Db	721	TCAGAGCTGAATACAGGCAATTTATTTCAACAGTACAGCAAGCACTGATGATGAGCACTTA	780
QY	928	ACAGGTCCCCCAAGCAAGAACTATTTATGCACTCAAGTTTACCAAGCTCAGCTGGCT	987
Db	781	ACAGGTCCCCCAAGCAAGAACTATTTCTCATGCAATCAAGTTTACCCAGGCTCAGCTGGCT	840
QY	988	TCATATGAATCTTCTGACATTTGATCAAGATGGAAGAACTTTACAGCAGAGGAATTTATC	1047
Db	841	TCATCTGGAATCTTTCCGACATTTGATCAAGATGGAAGAACTTTACCCAGCAAGAAATTTATC	900
QY	1048	CTGGCAATCACCTCATGATGATGTATGCTTGGCAACCACTGCACTGCTGCTGCT	1107
Db	901	CTAGCGATGACCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG	960
QY	1108	CCAGATATCATCTTCTTTTGAAGAGTTTCTGATCTGGCAGTGTATATCTGTCTATA	1167
Db	961	CCAGATATCATCTTCTTCTTTGAAGAGTTTCTGATCTGGCAGTGTATATCTGTCTATA	1020
QY	1168	AGCTCAACATCTGTAGATCAGAGGCTTACAGAGGAACCAAGTTTGAAGATGAACACAA	1227
Db	1021	AGCTCTTCTGTGACAGCAGCGGCTGCGCGAGGAGCCATCTGTCAGAGGATGAGCAGAG	1080
QY	1228	CAATAGAAAGAAATTTACTGTAACTTTTGAAGATAAGAAAGCGGAGAACTTTGAACCT	1287
Db	1081	G---TGGAAGAAAGCTGTGTGATTTTGAAGATAAGAAAGCGGAGAACTTTGAGCGA	1137
QY	1288	GGCAACTGTGGAACCGAAGGCAAGCTTCTCTGGAAACAGCAGCGCAAGGAGCAG	1347
Db	1138	GGCAACTGTGGAAGCGCAGCAGCGGCTCTCTGGAGCAGCAGCGCAAGGAGCAG	1197
QY	1348	GAGCGCTGGCCAGCTGAGCGGGCGGAGCAGGAGAGGAGGAGGAGGAGGAGGAGGAGGAG	1407
Db	1198	GAGCGCTGGCTCAGCTGGAGCGCGCAGAGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG	1257
QY	1408	CAAGAGCGCAAGAACTGGAACCTGGAAGCAACTGGAAGCAAGCGGAGCTTAGAA	1467
Db	1258	CAGGAGCGCAAGAGACAGCTAGAGCTGGAAGAGCAGCTTAGAAAGCGGAGCTGGAG	1317
QY	1468	CGGACAGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG	1527
Db	1318	AGGACAGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG	1377
QY	1528	CTTGAAGCGCAACGACAACTTTGATGGGAAACGGAATCGAAGCAAGAACTACTAAATCAA	1587
Db	1378	CTCGAAACGGCAGCGACAGCTTTGAATGGGAAACGGAATCGAAGCAAGAACTACTAAATCAA	1437
QY	1588	AGAAACAAAGAACCAAGAGGACATAGTTGTACTGTAAGAAACCAAGAAAGAACTTTTGAATTT	1647
Db	1438	AGGAACAAAGAACCAAGAGGAGGACATAGTTGTCTGTAAGAAACCAAGAAAGAACTTTTGAATTT	1497
QY	1648	GAATTAGAAGCTCTTAAATGATAAAGCATCAACTAGAGGGAACCTTCAAGATATCAGA	1707
Db	1498	GAATTAGAAGCTCTTAAATGATAAAGCATCAACTAGAGGGAACCTTCAAGATATCAGA	1557
QY	1708	TGTCGATTTGACACCCAAAGCAAGAAATTTAGAGCACAAACAAATCTAGAGAGTTGAGA	1767
Db	1558	TGTCGATTTGACACCCAAAGCAAGAAATTTAGAGCACAAACAAATCTAGAGAGTTGAGA	1617
QY	1768	ATTGCGGAAATCACCCATCTCAGCAACAAATTTACAGGAATCTCAGCAAAATGCTTGAAGA	1827
Db	1618	ATTGCGGAAATCACCCATCTCAGCAACAAATTTACAGGAATCTCAGCAAAATGCTTGAAGA	1677
QY	1828	CTTATTCAGAAAAACAGATCTCAATGACCAATTTAAACAAAGTTTCCAGCAGAACAGTTTG	1887
Db	1678	CTTATTCAGAAAAACAGATCTCAGTCAACCAATTTAAACAAAGTTTCCAGCAGAACAGTTTG	1737
QY	1888	CACAGAGTTCACTTTTACACTTAAAGAGCCTTTAGAGCAAAAGAACTAGATCAAAATCTACAGGAGTT	1947
Db	1738	CATAGAGATTTCTTCTTACCTTAAAGAGCCTTTGGAAGCAAAAGAACTAGATCAAAATCTACAGGAGTT	1797
QY	1948	CACCTTACAGAGCAAACTGGATGGAAGAAAGAACTTAGATCAAAATCTACAGGAGTT	2007
Db	1798	CAGCTTTCAGAGCAGCTGGACGAGTGGAGAAAGAGACAGCTCAAGCTCGCAGGAGATT	1857
QY	2008	GATATTTCAATATCAGCTGAAGGAACTAAGAGAAATACAAATTAAGCAAACTACCTCAG	2067
Db	1858	GATGTTTTTCAACACAGCTGAAGGAACTGAGAGAGATACACAGTAAACAGCAGCTCCAG	1917
QY	2068	AAAGCAAAAGTCCATGAGCTGAACTGAAAGCAAGAAAGAACTAGATCAAAATCTACAGGAGTT	2127
Db	1918	AAAGCAAAAGTCCATGAGCTGAACTGAAAGCAAGAAAGAACTAGATCAAAATCTACAGGAGTT	1977
QY	2128	GAATTTAGAAAAACAAAAAGAAAGCCCAAGACGAGCTCAGGAAAGGAAAGCAAGCTGG	2187
Db	1978	GAGTTGGAGAAAGCAAAAGAAAGGTCAGAGACGAGTTTCAAGAAAGGAAAGCAAGCAATGG	2037
QY	2188	CTGAGAGCTGTGACAGCAGAGGACAGCATCAGAGACCAAGAAACTCCACCAAGAGGAA	2247
Db	2038	CAGGAGCATGTCCAGCAGGAG---GAGCAGCAGCGCTTCCGAAACCCCAAGGAGGAC	2094
QY	2248	AAACTGAAAGGGGAGGAGTGTCAAAAGAAAGGATGGCAGGAAAGGCAAAAGGCAAGGAA	2307
Db	2095	AAACTGAAAGGGGAGGAGTGTCAAAAGAAAGGATGGCAGGAAAGGCAAAAGGCAAGGAA	2154
QY	2308	GCACAAGCAAGCTGGGCTTTCATCAACCAAGAAACCAAGCAAGCTTAAGCAGCAGCTGTC	2367
Db	2155	GTGCAAGCAAGCAGCAGCTGGCTTTTCCATCCACATCAAGACCCAGCTTAAGCCGGC---G	2211
QY	2368	CAGGCAACCTGTGTCTGCAAGAAAGGTCACCTTACCTTCTGCAAGGAAATGTA	2427
Db	2212	CAGGCAACCTGTGTCTGCAAGAAAGGTCACCTTACCTTCTGCAAGGAAATGTA	2271
QY	2428	AAAGTGGTGTATTATTCGGGCACTGTACCTTTTGAATCCAGAAAGGATGATGAATCACT	2487
Db	2272	AAAGTGGTGTATTATTCGGGCACTGTACCTTTTGAATCCAGAAAGGATGATGAATCACT	2331
QY	2488	ATCCAGCAGGAGACATAGTCAAT-----GGTGGATGAAGCCAAACTGGA	2532
Db	2332	ATCCAGCAGGAGACATAGTCAAT-----GGTGGATGAAGCCAAACTGGA	2391
QY	2533	GAAACCGGCTGTGAGGAGGAAATTAAGGAAAGACAGGCTGCTTCCCTCAGAACTAT	2592
Db	2392	GAGCCAGGATGCTTGGAGGAGAACCGAAAGGGGAGACAGGATGTTTCCCTCAGAACTAT	2451
QY	2593	GCAGAGAAAAATCCAGAAAAATGAGGTTCCCGCTCCAGTGAAACCAAGTGAATTTCAACA	2652

Db	130	GAGAAGTCAGTGGGCAGGTTCGGGCTCTTCCCGTCCAATTTATGTGAAGCTGCACACAGCA	189
Qy	3623	TGGACCCCAAGCCAGCAATGAATCATATGTTGTCCATCCCCCCTCAGGCTTGAAGTCCT	3682
Db	190	TGGACCCCAAGCCAGCAATGAATCATATGTTGTCCAT-CCCCCCTCAGGCTTGAAGCCCT	248
Qy	3683	CAAAAGAGACCCACTATCCCATATCACTGCCCCAGAGGGATGATGGGAGATGCAGCCTTGAT	3742
Db	249	CAAAAGAGACCCACTATCCCATATCACTGCCCCAGAGGGATGATGGGAGATGCAGCCTTGAT	308
Qy	3743	CATGTGAATTCAGCATGATCACTACTGCCCTTCTGAGTAGAAGAACTCACTCAGAGCA	3802
Db	309	CATGTGAATTCAGCATGATCACTACTGCCCTTCTGAGTAGAAGAACTCACTCAGAGCA	368
Qy	3803	GTTTACCTCATTTTACCTTAGTTGCATGTGTCGCAATCTTTTGAGTTATTTACCTGCAGAG	3862
Db	369	GTTTACCTCATTTTACCTTAGTTGCATGTGTCGCAATCTTTTGAG-TCCCTGGCTGCAGAG	427
Qy	3863	ATAGGAGCAAAAATTTACAAAAACACACAGGGTAGTGGGTCCCTTTTGTGGCTTTCCCTAGTT	3922
Db	428	GCAGAGC--AAATTGCAGAACTGCACAGGGTGGGGGTCCTTTTGT- GCTTTCCTAGTC	484
Qy	3923	ACTCAAAATTGACTTTTCCCCACCTTTTGCA CAGGTGCTTTTCAATAGTTTTTAAAAATTATTTT	3982
Db	485	ACTCAGACTGACGGCCCGCCCTTCASWCWAGVGACTWKSAA-AGYYWYAARAKAVTTT	543
Qy	3983	TAAATATATATTTTAGCTTTTAAATAACA	4012
Db	544	TAAATGTGTAATTTTAGCTTTTAAATAAAA	573

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RESULT 3
US-11-091-883-433
; Sequence 433, Application US/11091883
; Publication No. US20060024693A1
; GENERAL INFORMATION:
; APPLICANT: CIBELLI, JOSE
; APPLICANT: FERNANDEZ, EMILIO O.
; APPLICANT: JORDAO DE MEGALHAES, GUILHERME
; APPLICANT: KOCABAS, ARIF
; APPLICANT: CROSBY, JAVIER A.
; TITLE OF INVENTION: IDENTIFICATION OF GENES OR POLYPEPTIDES THE EXPRESSION OF WHICH
; TITLE OF INVENTION: CORRELATES TO FERTILITY, OVARIAN FUNCTION AND/OR FETAL/NEWBORN
; TITLE OF INVENTION: VIABILITY
; FILE REFERENCE: 53942US
; CURRENT APPLICATION NUMBER: US/11/091,883
; CURRENT FILING DATE: 2005-03-29
; PRIOR APPLICATION NUMBER: 60/556,875
; PRIOR FILING DATE: 2004-03-29
; NUMBER OF SEQ ID NOS: 513
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 433
; LENGTH: 4053
; TYPE: DNA
; ORGANISM: Homo sapiens
US-11-091-883-433

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Qy	3388	AGCCTGGGAGCAGCAAAATCACTCCAACAGAGCCACCTAAGTCAACAGACATTAGCGGCA	3447
Db	514	GGTCCAAGCAGTGAAGAGGCA-----CACCTGCTTTTCATCCT	552
Qy	3448	GTCTGCCAGGTGATTCGGGATGTACGACTACCCGCGCAGATCAGCATGAGCTGGCCCTTC	3507
Db	553	GTAATGTCAGGTGATTCGTAATGATGACTATGCAGCAAAATATGAAGATGAGCTCAGTTTC	612
Qy	3508	AACAAGGGCCAGATCATCAACGTCCTCAACAAGGAGGACCCCTGACTGGTGGAAAGAGAA	3567
Db	613	TCCAAGGGCAACTCATTAATGTTATGAAACAAGATGATCCTGATTGGTGGCAAGGAGAG	672
Qy	3568	GTCAATGGACAAGTGGGGCTCTCCCATCCAATTATGTGNAAGCTGACCACAGACATGGAC	3627
Db	673	ATCAACGGGGTGACTGGTCTCTTTCTCTTCAAACTAGCTTAAGATGACGACAGACTCAGAT	732
Qy	3628	CCAAGCCAGCAATGAATCATATGTTGTCATCCCCCTCAGCGCTTGAAGTCTCTCAAAG	3687
Db	733	CCAAGTCAACAGTGAACCA-----ATGTTGTCCTCCAGTGTGAAAGCACCCAG	782
Qy	3688	AGACCCACTATCCCATATCACTGCCAGAGGGATGATGGAGATGCGCCTTGATCATGT	3747
Db	783	AGACCCACTAT-CCAAGTTTCACTTAGCGTGGAGCAGGCGAGGCCCTGATCAAAAT	841
Qy	3748	GACTTCCAGCATGATCACCTACTGCCTT	3775
Db	842	ATCTCTACACAATCGTTTACTTCGTT	869

RESULT 4
 US-10-750-185-31648
 ; Sequence 31648, Application US/10750185
 ; Publication No. US20050260603A1
 ; GENERAL INFORMATION:
 ; APPLICANT: MMI GENOMICS, INC.
 ; APPLICANT: DENISE, Sue K.
 ; APPLICANT: KERR, Richard
 ; APPLICANT: ROSENFELD, David
 ; APPLICANT: HOLM, Tom
 ; APPLICANT: BATES, Stephen
 ; APPLICANT: FANTIN, Dennis
 ; TITLE OF INVENTION: COMPOSITIONS FOR INFERRING BOVINE TRAITS
 ; FILE REFERENCE: MW1100-2
 ; CURRENT APPLICATION NUMBER: US/10/750,185
 ; CURRENT FILING DATE: 2003-12-31
 ; PRIOR APPLICATION NUMBER: US 60/437,482
 ; PRIOR FILING DATE: 2002-12-31
 ; NUMBER OF SEQ ID NOS: 64922
 ; SOFTWARE: PatentIn version 3.1
 ; SEQ ID NO 31648
 ; LENGTH: 874
 ; TYPE: DNA
 ; ORGANISM: Bovine
 US-10-750-185-31648

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Db      281 ATCCCTGGG 289
|| | | |
Best Local Similarity 51.5%; Pred. No. 6.7e-06;
Matches 177; Conservative 0; Mismatches 167; Indels 0; Gaps 0;

RESULT 5
US-10-750-623-31648
; Sequence 31648, Application US/10750623
; Publication No. US20050287531A1
; GENERAL INFORMATION:
; APPLICANT: MMI GENOMICS, INC.
; APPLICANT: DENISE, Sue K.
; APPLICANT: KERR, Richard
; APPLICANT: ROSENFELD, David
; APPLICANT: HOLM, Tom
; APPLICANT: BATES, Stephen
; APPLICANT: FANTIN, Dennis
; TITLE OF INVENTION: METHODS AND SYSTEMS FOR INFERRING BOVINE TRAITS
; FILE REFERENCE: MM1100-1
; CURRENT APPLICATION NUMBER: US/10/750,623
; CURRENT FILING DATE: 2003-12-31
; PRIOR APPLICATION NUMBER: US 60/437,482
; PRIOR FILING DATE: 2002-12-31
; NUMBER OF SEQ ID NOS: 64922
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 31648
; LENGTH: 874
; TYPE: DNA
; ORGANISM: Bovine 19866880882217
US-10-750-623-31648

Query Match      1.9%; Score 96.2; DB 7; Length 874;
Best Local Similarity 69.3%; Pred. No. 4.2e-11;
Matches 131; Conservative 0; Mismatches 58; Indels 0; Gaps 0;

QY      2917 CAGGCTCAAAAGGTGGAGGGCTACAGCTCAAGCCCTATATCTCTGGAGAGCAAAAAA 2976
Db      101 CAGGACAGGTTGTAGAAAACCTGAAGCACAGGCCCTTTGTCTTGACTGCAAGAAA 160

QY      2977 GACAACACCTTAATATTTTAAACAAAATGATGTATCATCACCGTCTCTGGAAACAGCAGACATG 3036
Db      161 GAAAACCACTTGAACCTTCTCAAAACATGATATTATTACTGTCTTAGAGCAGCAAGAAAT 220

QY      3037 TGGTGGTTTGGAGAGTTCAAGTCAGAGGGTTGGTCCCAAGTCTTACGTGAACATC 3096
Db      221 TGGTGGTTTGGGAGGTTTCACGGAGGAAGGATGGTTCCGAAATCTTACGTCAAAATC 280

QY      3097 ATTTTCAGG 3105
Db      281 ATCCCTGGG 289

RESULT 6
US-11-121-086-5
; Sequence 5, Application US/11121086
; Publication No. US20050266459A1
; GENERAL INFORMATION:
; APPLICANT: POULSEN, TIM S.
; APPLICANT: NIELSEN, KIRSTEN V.
; TITLE OF INVENTION: NUCLEIC ACID PROBES AND NUCLEIC ACID ANALOG PROBES
; FILE REFERENCE: 09138.6000-00000
; CURRENT APPLICATION NUMBER: US/11/121,086
; CURRENT FILING DATE: 2005-05-04
; PRIOR APPLICATION NUMBER: 60/567,570
; PRIOR FILING DATE: 2004-05-04
; NUMBER OF SEQ ID NOS: 107
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 5
; LENGTH: 153376
; TYPE: DNA
; ORGANISM: Homo sapiens
US-11-121-086-5

Query Match      1.5%; Score 76.8; DB 11; Length 153376;

Best Local Similarity 51.5%; Pred. No. 6.7e-06;
Matches 177; Conservative 0; Mismatches 167; Indels 0; Gaps 0;

QY      1183 GATCAGAGGCTACCAGAGAACCCAGTTTGTAGAGATGAACAACAATTTAGNAAGAAA 1242
Db      15959 GAGGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 16018

QY      1243 TTACCTGTAACTTTGAAGATAAGAGCGGAGAGACTTTTGAACGTGGCAACCTGGAACTG 1302
Db      16019 TAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 16078

QY      1303 GAGAAACGAAAGCAAGCTCTCTCTGGAACAGCAGCGGCAAGGAGGAGGAGGAGGAGGAGGAG 1362
Db      16079 GAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 16138

QY      1363 CTGGAGCGGCGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1422
Db      16139 GAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 16198

QY      1423 CAACTGGAACCTGGAGAAAGCAACTGGAAAGAGCAGCGGAGCTAGAACGGCAGAGAGGAGGAG 1482
Db      16199 GAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 16258

QY      1483 GAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1526
Db      16259 GAACAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGA 16302

RESULT 7
US-10-750-185-31653
; Sequence 31653, Application US/10750185
; Publication No. US20050260603A1
; GENERAL INFORMATION:
; APPLICANT: MMI GENOMICS, INC.
; APPLICANT: DENISE, Sue K.
; APPLICANT: KERR, Richard
; APPLICANT: ROSENFELD, David
; APPLICANT: HOLM, Tom
; APPLICANT: BATES, Stephen
; APPLICANT: FANTIN, Dennis
; TITLE OF INVENTION: COMPOSITIONS FOR INFERRING BOVINE TRAITS
; FILE REFERENCE: MM1100-2
; CURRENT APPLICATION NUMBER: US/10/750,185
; CURRENT FILING DATE: 2003-12-31
; PRIOR APPLICATION NUMBER: US 60/437,482
; PRIOR FILING DATE: 2002-12-31
; NUMBER OF SEQ ID NOS: 64922
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 31653
; LENGTH: 2170
; TYPE: DNA
; ORGANISM: Bovine 19866880980841
US-10-750-185-31653

Query Match      1.4%; Score 74; DB 7; Length 2170;
Best Local Similarity 67.5%; Pred. No. 3.9e-06;
Matches 104; Conservative 0; Mismatches 50; Indels 0; Gaps 0;

QY      986 CTTCAATATGGAATCTTTCTGCAATTCATGATCAAGATGAAAACTTACAGCAGAGGAATTA 1045
Db      1466 CTTCTTTCAGGACTCTGGCTGACATCGATCGTGTATGGAACAGCTGAAAGCTGAGGAGTTCA 1525

QY      1046 TCCTGGCAATGCACCTCATTGATGTAGCTATGTCTGCCCAACCACTGCCACCTCTCTCTGC 1105
Db      1526 TCCTTGGATGACCTCCTACTGATATGATGATGATGATGATGATGATGATGATGATGATGAT 1585

QY      1106 CTCAGAAATACATTTCCACCTTTCTTTTGAAGAGT 1139
Db      1586 CTCCTGAGCTTGCTCCCTCCATCTTTTAGGTGAGT 1619

RESULT 8
US-10-750-623-31653
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; CURRENT APPLICATION NUMBER: US/11/136,527
; CURRENT FILING DATE: 2005-05-25
; PRIOR APPLICATION NUMBER: US 60/574,294
; PRIOR FILING DATE: 2005-05-26
; NUMBER OF SEQ ID NOS: 362830
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 3376
; LENGTH: 2380
; TYPE: DNA
; ORGANISM: Rattus norvegicus
US-11-136-527-3376

Query Match      1.2%; Score 64.2; DB 11; Length 2380;
Best Local Similarity 55.1%; Pred. No. 0.00052;
Matches 147; Conservative 0; Mismatches 118; Indels 2; Gaps 1;

QY 2066 AGAAGCAAAAGTCCATGGAGGCTGAACGACTGAAACACAGAAACCAAGAACGAAATCA 2125
DB 1512 AGAGGAAGAGAGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 1453
QY 2126 TAGAATTAGAAAAACAAAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 2185
DB 1452 AAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 1395
QY 2186 GCGTGGAGCAGTGTGACGAGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 2245
DB 1394 AGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAG 1335
QY 2246 AAAAAGTGAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 2305
DB 1334 AGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 1275
QY 2306 AAGCACAAGACAAGCTGGGTGGCTTT 2332
DB 1274 AAGAGAGAGTCCAGCAGTGTGGTCTTT 1248

RESULT 12
US-11-136-527-155
; Sequence 155, Application US/11/136527
; Publication No. US20050287570A1
; GENERAL INFORMATION:
; APPLICANT: Wyeth
; TITLE OF INVENTION: Mounts, William M
; FILE REFERENCE: 031896-041000 (AM101086)
; CURRENT APPLICATION NUMBER: US/11/136,527
; CURRENT FILING DATE: 2005-05-25
; PRIOR APPLICATION NUMBER: US 60/574,294
; PRIOR FILING DATE: 2005-05-26
; NUMBER OF SEQ ID NOS: 362830
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 155
; LENGTH: 3456
; TYPE: DNA
; ORGANISM: Rattus norvegicus
US-11-136-527-155

Query Match      1.2%; Score 63.8; DB 11; Length 3456;
Best Local Similarity 58.6%; Pred. No. 0.00075;
Matches 99; Conservative 6; Mismatches 64; Indels 0; Gaps 0;

QY 3439 TTAGCGCAGTGTGCCAGGTGATGGATGATGACGACTACCGCGCAGAGATGACGATGAG 3498
DB 434 TTGSCDCAATGGTGGAGGCCATAGTGGAGTTTGATTACCAGGCCAGCATGATGATGAG 493
QY 3499 CTGCGCTTCAACAGAGGCCAGATCATCACTCTCTCAACAGAGGAGGCCCTGACTGTGG 3558
DB 494 CTGACGATCAGCGTGGGTGAGGTGATCATCACWACWCTCAGWAGWAGGATGGAGGCTGGTGG 553
QY 3559 AAAGGAGAGTCAATGGACAGTGGGCTCTTCCCATCCATTTATGTGA 3607
DB 554 GAGGACAGATCAACGGCAGGAGAGGTTTGTTCCTGACACTTTGTAA 602
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RESULT 13
US-11-011-332A-156/c
; Sequence 156, Application US/11011332A
; Publication No. US20060024684A1
; GENERAL INFORMATION:
; APPLICANT: Roekens, John
; APPLICANT: Harbeck, Nadia
; APPLICANT: Koenig, Thomas
; APPLICANT: Maier, Sabine
; APPLICANT: Martens, John
; APPLICANT: Model, Fabian
; APPLICANT: Nimmrich, Inko
; APPLICANT: Rujan, Tamas
; APPLICANT: Schmitt, Manfred
; APPLICANT: Lesche, Ralf
; APPLICANT: Dietrich, Dima
; APPLICANT: Mueller, Volkmar
; APPLICANT: Kluth, Antje
; APPLICANT: Schweppe, Ina
; APPLICANT: Hartmann, Oliver
; APPLICANT: Adorjan, Peter
; TITLE OF INVENTION: PROGNOSTIC MARKERS FOR PREDICTION OF TREATMENT RESPONSE AND/OR S
; FILE REFERENCE: 47675-99
; CURRENT APPLICATION NUMBER: US/11/011,332A
; CURRENT FILING DATE: 2004-12-13
; PRIOR APPLICATION NUMBER: US 10/517,741
; PRIOR FILING DATE: 2003-10-01
; PRIOR APPLICATION NUMBER: PCT/EP2003/010881
; PRIOR FILING DATE: 2003-10-01
; PRIOR APPLICATION NUMBER: DE 10245779.4
; PRIOR FILING DATE: 2002-10-01
; PRIOR APPLICATION NUMBER: DE 10300096.8
; PRIOR FILING DATE: 2003-01-07
; PRIOR APPLICATION NUMBER: DE 10317955.0
; PRIOR FILING DATE: 2003-04-17
; PRIOR APPLICATION NUMBER: PCT/EP2004/014170
; PRIOR FILING DATE: 2004-12-13
; PRIOR APPLICATION NUMBER: EP 03090432.0
; PRIOR FILING DATE: 2003-12-11
; PRIOR APPLICATION NUMBER: EP 04090041.7
; PRIOR FILING DATE: 2004-02-10
; PRIOR APPLICATION NUMBER: EP 04090380.9
; PRIOR FILING DATE: 2004-09-30
; PRIOR APPLICATION NUMBER: EP 04090127.4
; PRIOR FILING DATE: 2004-04-01
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 158
; SEQ ID NO 156
; LENGTH: 28536
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
US-11-011-332A-156

Query Match      1.2%; Score 63.8; DB 11; Length 28536;
Best Local Similarity 45.7%; Pred. No. 0.002;
Matches 273; Conservative 0; Mismatches 312; Indels 12; Gaps 1;

QY 1561 AATCGAGGCGAAGCACTACTTAATCAAGAAACAAAGAAAGAGGACATAGTTGTACTG 1620
DB 6698 AATCAAAAAACAAACCAAAAAACAAAAACAAAAACAACTTAACTTAACTTAACTTAA 6639
QY 1621 AAAGCAAGAAAAAGACTTTTGGATTTGAATTTGAAGCTCTTAAATGATAAAAGCATCAA 1680
DB 6638 CAACCAAAAAACAAACACACAACTCCCACTATATAAAAAACCAACCAAAAAACATAA 6579
QY 1681 CTAGAGGGAACTTCAAGATATATGATGTGATGTGATGTGATGTGATGTGATGTGATG 1740
DB 6578 AATAAAAAACAAAAACAACTTACACCAACCAACCAACCTTAATTAATTAATTAATTA 6519
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1741 AGCACAAACAAATCTAGAGATTGAGAAATTGCGGAAATCACCCATCTACAGCAACAATTA 1800
| | | | |
6518 AACACCAAAACAAACAAAAACAAAAACATACTTTTAAAAAACAATCAAAACAAACTAA 6459
| | | | |
1801 CAGGAATCTCAGCAATGCTTGAAGACTTATTCAGAAAAACAGATCTCAATGACCAA 1860
| | | | |
6458 AAAAAACAAACAAAAAATAACCCCAAAAAACAAACAAAAACAAATATTCTATAAATTA 6399
| | | | |
1861 TTAATAACAGATTGAGCAGAACAGTTTGCACAGAGATTCACTTGTACACTTAAAGAGCC 1920
| | | | |
6398 TTTAAACAAACAAATATAAATAATAATAAATACATATATAACCTTCTTCAACCC 6339
| | | | |
1921 TTAGAAGCAAAAGAACTAGCTCGGCAGCACTACGAGACCAACTGGATGAAGTGGAGAA 1980
| | | | |
6338 TACAAAATTAATAAAAAAATACAAACAAAAAATACTACAAA-----AAAA 6291
| | | | |
1981 GAACTAGATCAAAACTACAGAGATTGATATTTTCAATAATCAGCTGAAGCACTAAGA 2040
| | | | |
6290 AAAAAATAAAAAACCAATAAAAAACAAAAATNAAAATTAACAAATAAATAAATAAAAA 6231
| | | | |
2041 GAAATACACAATAAGCAACAACTCCAGAAGCAAAAGTCCATGAGGCTGAAGCACTGAA 2100
| | | | |
6230 TAAATAAAAAAATCTAAAAAATAAATAAATAAATAAATAAATAAATAAATAAATAA 6171
| | | | |
2101 CAGAAAGAAACAAAGAAAGATCATAGAAATTAGAAAAACAAAAAGAAAGCCCAA 2157
| | | | |
6170 ACCAAAAACAAACAACTAACTTATCAACTAAAAAATAAATAAATAAATAAATAAATAA 6114
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RESULT 14

US-11-000-688-163
; Sequence 163, Application US/11000688
; Publication No. US20050287544A1
; GENERAL INFORMATION:
; APPLICANT: BERTUCCI, Francois
; APPLICANT: HOULGATTE, Remi
; APPLICANT: BIRNBAUM, Daniel
; TITLE OF INVENTION: GENE EXPRESSION PROFILING OF COLON CANCER WITH DNA ARRAYS
; FILE REFERENCE: 1423-R-03
; CURRENT APPLICATION NUMBER: US/11/000,688
; CURRENT FILING DATE: 2004-12-01
; PRIOR APPLICATION NUMBER: US 60/525,987
; PRIOR FILING DATE: 2003-12-01
; NUMBER OF SEQ ID NOS: 1596
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 163
; LENGTH: 4257
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial sequences:primer
; NAME/KEY: misc feature
; LOCATION: (1)..(4257)
; OTHER INFORMATION: epidermal growth factor receptor pathway
; OTHER INFORMATION: substrate 15(BPS15) gene.
US-11-000-688-163

Query Match 1.2%; Score 63.2; DB 11; Length 4257;
Best Local Similarity 56.9%; Pred. No. 0.0011;
Matches 116; Conservative 0; Mismatches 88; Indels 0; Gaps 0;

250 TGGGCCATAACTGTAGAGAAAGAGCGAAGCATGTATCAGCAGTTCATAGTTTAAAGCCA 309
| | | | |
456 TGGGCTGTAAACCTGAAGTAAGGCCAAATATGATGCAATATTGATAGTTTAAAGCCA 515
| | | | |
310 ATATCTGGATTCTACTCTGGTATCAAGCTAGAAACCTTTTTTTTCAATCTGGGTACCT 369
| | | | |
516 GTGAATGGAAATTTCTGTCTGGTATGAAGTGAACCCAGTGTGCTCAACTCTAAGTTACCT 575
| | | | |
370 CAACCTGTTTATAGCAGAGATATGGCACTAGCTGACATGAATAATGATGAAGAAATGGAT 429
| | | | |

576 GTGGATATCCTTTGGAGAGATTGGAGTTGAGTATATTGACCATGATGGAATGCTTGAC 635
| | | | |
430 CAAAGTGGAGTTTTCATAGCTATG 453
| | | | |
636 AGAGATGAGTTTGACAGTTGCCATG 659
| | | | |
RESULT 15
US-11-000-688-1278
; Sequence 1278, Application US/11000688
; Publication No. US20050287544A1
; GENERAL INFORMATION:
; APPLICANT: BERTUCCI, Francois
; APPLICANT: HOULGATTE, Remi
; APPLICANT: BIRNBAUM, Daniel
; TITLE OF INVENTION: GENE EXPRESSION PROFILING OF COLON CANCER WITH DNA ARRAYS
; FILE REFERENCE: 1423-R-03
; CURRENT APPLICATION NUMBER: US/11/000,688
; CURRENT FILING DATE: 2004-12-01
; PRIOR APPLICATION NUMBER: US 60/525,987
; PRIOR FILING DATE: 2003-12-01
; NUMBER OF SEQ ID NOS: 1596
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 1278
; LENGTH: 2774
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial sequences:primer
; NAME/KEY: misc feature
; LOCATION: (1)..(2774)
; OTHER INFORMATION: epidermal growth factor receptor substrate
; OTHER INFORMATION: eps15r(BPS15R) gene.
US-11-000-688-1278

Query Match 1.2%; Score 62.8; DB 11; Length 2774;
Best Local Similarity 52.3%; Pred. No. 0.0011;
Matches 139; Conservative 0; Mismatches 127; Indels 0; Gaps 0;

249 CTGGGCCATAACTGTAGAGAAAGAGCGAAGCATGTATCAGCAGTTCATAGTTTAAAGCC 308
| | | | |
366 CTGGGCTGTAGGCTGGAGAAAGGCCAAATTTGATGGGATTTTGAAGCCTCTTGCC 425
| | | | |
309 AATATCTGGATTCTACTCTGGTATCAAGCTAGAAACCTTTTTTTTCAATCTGGGTACC 368
| | | | |
426 CATCAATGGTTTGCTCTCTGGAGACAAAGTCAAGCCAGTCTCTCATGAACCTCAAGCTGCC 485
| | | | |
369 TCAACCTGTTTATAGCAGAGATATGGCACTAGCTGACATGAATATATGATGAAGATGGA 428
| | | | |
486 TCTTGATGTCTGGGACGGTCTGGACCTCAGTGACATTGACAAGGATGGGCACTTGA 545
| | | | |
429 CAAAGTGGAGTTTTCATAGCTATGAAACTTATCAAACTGAAGCTTACAAGGATATCAGCT 488
| | | | |
546 TCGAGATGAGTTCGCTGTGGCCATGCACTTGGTGTACCGAGCCCTGGAGAGAGCCCT 605
| | | | |
489 ACCCTCTGCACCTTCCCTCTGTATGA 514
| | | | |
606 GCCCTCCGCTGCGCCGCTCCCTCA 631
| | | | |

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Job time : 752 secs

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OM nucleic - nucleic search, using sw model

Run on: February 14, 2006, 05:22:49 ; Search time 3772 Seconds
(without alignments)
11397.799 Million cell updates/sec

Title: US-09-720-934-1
Perfect score: 5199
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Scoring table: IDENTITY NUC
Gapop 10.0 , Gapext 1.0

Searched: 9793542 seqs, 4134689005 residues
Total number of hits satisfying chosen parameters: 19587084

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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8: /cgn2_6/ptodata/1/pubpna/US10D_PUBCOMB.seq:*
9: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*
10: /cgn2_6/ptodata/1/pubpna/US11_PUBCOMB.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	DB ID	Description
1	3285.6	63.2	7247	8	US-10-852-943-84
2	3285.6	63.2	7247	9	US-10-287-436A-57
3	3285.6	63.2	7247	9	US-10-287-436A-685
4	3090.6	59.4	7435	9	US-10-450-763-20567
5	2884.8	55.5	3466	7	US-10-158-057-33
6	2843.2	54.7	3319	6	US-09-764-875-88
7	1882.4	36.2	2067	6	US-10-264-049-887
8	1630	31.4	2874	9	US-10-450-763-20566
9	550	10.6	5828	7	US-10-398-885A-15
10	507.8	9.8	2017	3	US-09-884-441-72
11	507.8	9.8	2017	3	US-09-907-969-72
12	507.8	9.8	2017	3	US-09-827-271-72
13	507.8	9.8	2017	6	US-10-198-053-72
14	507.8	9.8	2017	6	US-10-860-790-72
15	503.6	9.7	568	3	US-09-764-881-55
16	503.6	9.7	568	3	US-09-764-881-55
17	503.6	9.7	568	3	US-09-764-875-404
18	503.6	9.7	568	6	US-10-242-747-55
19	503.6	9.7	568	7	US-10-158-057-127
20	462.6	8.9	503	3	US-09-918-995-31258
21	291.4	5.6	395	9	US-10-450-763-20563
22	264.2	5.1	301	3	US-09-864-761-17146
23	263	5.1	263	3	US-09-864-761-17644, A

24	250	4.8	270	3	US-09-864-761-17127	Sequence 17127, A
25	250	4.8	286	3	US-09-864-761-26948	Sequence 26948, A
26	250	4.8	297	3	US-09-864-761-30453	Sequence 30453, A
27	240	4.6	486	3	US-09-864-761-333	Sequence 333, App
28	240	4.6	487	3	US-09-864-761-864	Sequence 864, App
29	239	4.6	480	3	US-09-864-761-10314	Sequence 10314, A
30	237.8	4.6	418	3	US-09-783-590-8760	Sequence 8760, App
31	230	4.4	230	3	US-09-864-761-17643	Sequence 17643, A
32	230	4.4	247	3	US-09-864-761-30501	Sequence 30501, A
33	205	3.9	484	3	US-09-864-761-863	Sequence 863, App
34	202	3.9	475	3	US-09-864-761-311	Sequence 311, App
35	202	3.9	475	3	US-09-864-761-13884	Sequence 13884, A
36	186	3.6	477	3	US-09-864-761-13936	Sequence 13936, A
37	180	3.5	180	3	US-09-864-761-17125	Sequence 17125, A
38	167.6	3.2	3981	10	US-11-097-143-233	Sequence 233, App
39	165.6	3.2	967	3	US-09-764-881-50	Sequence 50, Appl
40	165.6	3.2	967	3	US-09-764-881-50	Sequence 50, Appl
41	165.6	3.2	967	3	US-09-764-875-411	Sequence 411, App
42	165.6	3.2	967	6	US-10-242-747-50	Sequence 50, Appl
43	165.6	3.2	967	7	US-10-158-057-40	Sequence 40, Appl
44	165.4	3.2	292	3	US-09-864-761-20261	Sequence 20261, A
45	165.4	3.2	304	3	US-09-864-761-21373	Sequence 21373, A

ALIGNMENTS

RESULT 1
US-10-852-943-84
; Sequence 84, Application US/10852943
; Publication No. US20050037388A1
; GENERAL INFORMATION:
; APPLICANT: University of Geneva
; APPLICANT: Stylianos, Antonarakis
; APPLICANT: Deutsch, Samuel
; TITLE OF INVENTION: METHOD FOR DETECTING DISEASES CAUSED BY CHROMOSOMAL IMBALANCES
; FILE REFERENCE: 27067/2005
; CURRENT APPLICATION NUMBER: US/10/852,943
; CURRENT FILING DATE: 2004-05-25
; PRIOR APPLICATION NUMBER: US 60/300,266
; PRIOR FILING DATE: 2001-06-21
; PRIOR APPLICATION NUMBER: US 10/177,063
; PRIOR FILING DATE: 2002-06-21
; NUMBER OF SEQ ID NOS: 98
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 84
; LENGTH: 7247
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-852-943-84

Query Match	63.2%	Score	3285.6	DB	8	Length	7247
Best Local Similarity	93.8%	Pred. No.	0				
Matches	3536	Conservative	0	Mismatches	4	Indels	228
						Gaps	2

Qy	102	GCCTCCCTCCACGGCGCTGAGCGCACTGATTTCCTCGGGCGGCGAGCCGCGACC	161
Db	1	CGCTCCCTCCACGGCGCTGAGCGCACTGATTTCCTCGGGCGGCGAGCCGCGACC	60
Qy	162	CGCCCGGAGATGAGCGCTCGATTAGCAAGTAAAGTAAACAGAACCATGGCTCAGTTTCC	221
Db	61	CGCCCGGAGATGAGCGCTCGATTAGCAAGTAAAGTAAACAGAACCATGGCTCAGTTTCC	120
Qy	222	AACACCTTTGGTGGCGACCTCGATATCTGGCCATACCTGTAGAGAAAGAGCGAGCA	281
Db	121	AACACCTTTGGTGGCGACCTCGATATCTGGCCATACCTGTAGAGAAAGAGCGAGCA	180
Qy	282	TGATCAGCAGTTCATAGTTTAAAGCCATATCTGGATTCTGATTCATGATCAAGCTAG	341
Db	181	TGATCAGCAGTTCATAGTTTAAAGCCATATCTGGATTCTGATTCATGATCAAGCTAG	240
Qy	342	AAACCTTTTTCATCTCGGTTACTCAACCTGTTTATAGCAGATATGGGCACCTAGC	401

Db 241 AAACCTTTTTTTTCAATCTGGGTTACCTCAACCTGTTTTAGCACAGATATGGGCACTAGC 300
Qy 402 TGACATGAATAATGATGGAAGAATGGATCAAGTGGAGTTTTTCCATAGCTATGAAACTTAT 461
Db 301 TGACATGAATAATGATGGAAGAATGGATCAAGTGGAGTTTTTCCATAGCTATGAAACTTAT 360
Qy 462 CAAACTGAAGCTACAAAGGATATCAGCTACCTCTGCACTTCCCTCTGTCATGAAACAGCA 521
Db 361 CAAACTGAAGCTACAAAGGATATCAGCTACCTCTGCACTTCCCTCTGTCATGAAACAGCA 420
Qy 522 ACCAGTTGCTATTTCTAGCGCACCAAGCATTTGGTATGGAGGTATGCCAGCATGCCACC 581
Db 421 ACCAGTTGCTATTTCTAGCGCACCAAGCATTTGGTATGGAGGTATGCCAGCATGCCACC 480
Qy 582 GCTTACAGCTGTGCTCCAGTCCCAATGGGATCCATTCAGTTGTTGGAATGCTCCAAC 641
Db 481 GCTTACAGCTGTGCTCCAGTCCCAATGGGATCCATTCAGTTGTTGGAATGCTCCAAC 540
Qy 642 CCTAGTATCTTCTGTTCCCAAGCAGCTGTGCCCTCCCTGGCTTAAGGGGCTCCCTCTGT 701
Db 541 CCTAGTATCTTCTGTTCCCAAGCAGCTGTGCCCTCCCTGGCTTAAGGGGCTCCCTCTGT 600
Qy 702 TATACAACTCTGCTGCAATTTGCTCATCTGCAGCCACATTTGCCAAAGAGTTCTTCTTT 761
Db 601 TATACAACTCTGCTGCAATTTGCTCATCTGCAGCCACATTTGCCAAAGAGTTCTTCTTT 660
Qy 762 TAGTAGATCTGTTCCAGGCTCAACAATAAACACTAAATTAACAAAGGCAAGTCAATTTGA 821
Db 661 TAGTAGATCTGTTCCAGGCTCAACAATAAACACTAAATTAACAAAGGCAAGTCAATTTGA 720
Qy 822 TGTGGCAGTGTCCCAAGCTGGCAGAGTGGGCTGTTCTCAGTCATCAAGACTGAATA 881
Db 721 TGTGGCAGTGTCCCAAGCTGGCAGAGTGGGCTGTTCTCAGTCATCAAGCTGAATA 780
Qy 882 CAGGCAATTTATTCATAGTCATGACAAACTATAGTGGACACTTAAACAGTCCCAAGC 941
Db 781 CAGGCAATTTATTCATAGTCATGACAAACTATAGTGGACACTTAAACAGTCCCAAGC 840
Qy 942 AAGAACTATTTTATGCACTCAAGTTTACCACAGGCTCAGCTGGCTTCAATATGGAATCT 1001
Db 841 AAGAACTATTTTATGCACTCAAGTTTACCACAGGCTCAGCTGGCTTCAATATGGAATCT 900
Qy 1002 TTCTGACATGTACAGATGGAAGAACTTACAGCAGAGGAATTTATCTTGGCAATGCACCT 1061
Db 901 TTCTGACATGTACAGATGGAAGAACTTACAGCAGAGGAATTTATCTTGGCAATGCACCT 960
Qy 1062 CATTCATGTAGCTATGCTGGCCCAACCACTGCCACCTGCTCTGCTCCAGAAATACATTC 1121
Db 961 CATTCATGTAGCTATGCTGGCCCAACCACTGCCACCTGCTCTGCTCCAGAAATACATTC 1020
Qy 1122 ACCTTCTTTTGAAGAGTTTCAATCTGGCAGTGGTATATCTGTATAAGCTCAACATCTGT 1181
Db 1021 ACCTTCTTTTGAAGAGTTTCAATCTGGCAGTGGTATATCTGTATAAGCTCAACATCTGT 1080
Qy 1182 AGATCAGAGCTACCAAGCAACCACTTTTAGAAGTGAACCAACAAATTTAGAAGAA 1241
Db 1081 AGATCAGAGCTACCAAGCAACCACTTTTAGAAGTGAACCAACAAATTTAGAAGAA 1140
Qy 1242 ATTTACTGTAAAGTGTGAAGATAAGCGGAGAACTTTGAACCTGGCAACTGGAACT 1301
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Qy 1302 GGAGAAAAGCAAGCAAGCTCTCTCGAAACAGCAGCGCAAGGAGAGGCGCTGGCCCA 1361
Db 1201 GGAGAAAAGCAAGCAAGCTCTCTCGAAACAGCAGCGCAAGGAGAGGCGCTGGCCCA 1260
Qy 1362 GCTGAGCGGGCGGAGCAGAGAGGAGAGCGTGTAGCGCCAGGAGCAAGCGCAAGAG 1421
Db 1261 GCTGAGCGGGCGGAGCAGAGAGGAGAGCGTGTAGCGCCAGGAGCAAGCGCAAGAG 1320
Qy 1422 ACAACTGGAACTGGAGAACCAACTGGAAAAGCAGCGGGAGCTAGAACCGCAGAGAGGA 1481
Db 1321 ACAACTGGAACTGGAGAACCAACTGGAAAAGCAGCGGGAGCTAGAACCGCAGAGAGGA 1380

Qy 1482 GGAGAGGAGGAAAGAAATTTGAGCGGAGAGGCTGCAAAAACGGGAACCTTGAAGGCAACG 1541
Db 1381 GGAGAGGAGGAAAGAAATTTGAGCGGAGAGGCTGCAAAAACGGGAACCTTGAAGGCAACG 1440
Qy 1542 ACAACTTCAGTGGGAACGGAATCGAAGCAAGAACTACTAAATCAAGAAACAAAGAAACA 1601
Db 1441 ACAACTTCAGTGGGAACGGAATCGAAGCAAGAACTACTAAATCAAGAAACAAAGAAACA 1500
Qy 1602 AGAGGACATAGTTGTACTGAAAGCAAAAGAACTTTTGGAAATTTGAATTAGAAGCTCT 1661
Db 1501 AGAGGACATAGTTGTACTGAAAGCAAAAGAAAGACTTTTGGAAATTTGAATTAGAAGCTCT 1560
Qy 1662 AAATGATATAAAGCMTCAACTAGAAGGAAACTTCAAGATATCAGATCTCGATTCAGCCAC 1721
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Qy 1722 CCAAGGCAAGAAATTTGAGAGCACAAACAAATCTAGAGAGTTGAGAAATTCGCGAAATCAC 1781
Db 1621 CCAAGGCAAGAAATTTGAGAGCACAAACAAATCTAGAGAGTTGAGAAATTCGCGAAATCAC 1680
Qy 1782 CCATCTCAGCAACAATTTACAGGAATCTCAGCAAAATGCTTGGAGACTTATTTCCAGAAA 1841
Db 1681 CCATCTCAGCAACAATTTACAGGAATCTCAGCAAAATGCTTGGAGACTTATTTCCAGAAA 1740
Qy 1842 ACAGATCTCAATGACCAATTTAAACAAGTTTCAGCAGAACAGTTTGGCAGAGATTCAT 1901
Db 1741 ACAGATCTCAATGACCAATTTAAACAAGTTTCAGCAGAACAGTTTGGCAGAGATTCAT 1800
Qy 1902 TGTTCACCTTAAAGAGCTTTAGAGCAAAAGAACTAGCTCGGAGCACTTACAGAGCA 1961
Db 1801 TGTTCACCTTAAAGAGCTTTAGAGCAAAAGAACTAGCTCGGAGCACTTACAGAGCA 1860
Qy 1962 ACTGATGAAGTGGAGAAAGAACTTAGATCAAAACTACAGAGATTTGATATTTTCAATAA 2021
Db 1861 ACTGATGAAGTGGAGAAAGAACTTAGATCAAAACTACAGAGATTTGATATTTTCAATAA 1920
Qy 2022 TCAGCTGAAGAACTTAAGAGAAATACAAATAAGCAACAATCCAGAGCAAAAGTCCAT 2081
Db 1921 TCAGCTGAAGAACTTAAGAGAAATACAAATAAGCAACAATCCAGAGCAAAAGTCCAT 1980
Qy 2082 GGAGGCTCAACGACTGAAACAGAAAGAAACAAAGAACTATAGAAATTTAGAAAACA 2141
Db 1981 GGAGGCTCAACGACTGAAACAGAAAGAAACAAAGAACTATAGAAATTTAGAAAACA 2040
Qy 2142 AAAAGAAAGCCCAAGACGAGCTCAGGAAGGGAACGAGTGGCTGGAGCATGTGCA 2201
Db 2041 AAAAGAAAGCCCAAGACGAGCTCAGGAAGGGAACGAGTGGCTGGAGCATGTGCA 2100
Qy 2202 GCAGGAGCAGCATCAGAGACCAAGAAACTCCACCAAGGGAACCACTGAAAAGGGA 2261
Db 2101 GCAGGAGCAGCATCAGAGACCAAGAAACTCCACCAAGGGAACCACTGAAAAGGGA 2160
Qy 2262 GGAGAGTCTCAAAAAGAGGATGGCGAGAAAAGGCAACAGGAAGCACAAGACAAGCT 2321
Db 2161 GGAGAGTCTCAAAAAGAGGATGGCGAGAAAAGGCAACAGGAAGCACAAGACAAGCT 2220
Qy 2322 GGGTCGGCTTTTCCATCAACCAAGAACAGCTTAAGCCAGCTGTCCAGGCACTTGGTC 2381
Db 2221 GGGTCGGCTTTTCCATCAACCAAGAACAGCTTAAGCCAGCTGTCCAGGCACTTGGTC 2280
Qy 2382 CACTGCAGAAAAGGTCACCTTACCTTTCGCAAGGAAATGTAAAGTGGTCTATTA 2441
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Qy 2502 CATAGTCAT-----GGTGGATGAAAGCCAAACTGAGAGAACCCCGCTGGCT 2546
Db 2401 CATAGTCATGTTAAAGGGGAATGGTGGATGAAAGCCAAACTGAGAGAACCCCGCTGGCT 2460

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DB	481	GCTTACAGCTGTGCTCCAGTGGCAATGGATCCATTCAGTGTGTTGGAATGTCTCCAAC	540
QY	642	CCTAGTATCTTCTGTTCCACAGCAGCTGTGCCCCCTGGCTTAACGGGGCTCCCCCTGT	701
DB	541	CCTAGTATCTTCTGTTCCACAGCAGCTGTGCCCCCTGGCTTAACGGGGCTCCCCCTGT	600
QY	702	TATACAACTCTGCTGCTGCAATTTGCTCATCTCAGCCACATTTGCCAAGAGTTCCTTCCT	761
DB	601	TATACAACTCTGCTGCTGCAATTTGCTCATCTCAGCCACATTTGCCAAGAGTTCCTTCCT	660
QY	762	TAGTAGATCTGCTCAGGGTCACTAACTAAACACTAAATTTACAAAAGGCACAGTCATTGA	821
DB	661	TAGTAGATCTGCTCAGGGTCACTAACTAAACACTAAATTTACAAAAGGCACAGTCATTGA	720
QY	822	TGTGGCCAGTGTCCCAACAGTGGCAGTGGGCTGTTCTCAGTCATCAAGACTGAATA	881
DB	721	TGTGGCCAGTGTCCCAACAGTGGCAGTGGGCTGTTCTCAGTCATCAAGGCTGAATA	780
QY	882	CAGGCAATTAATCAATPAGTCATGACAAAACTATGAGTGGACACTTAAACAGGTCCTCAAGC	941
DB	781	CAGGCAATTAATCAATPAGTCATGACAAAACTATGAGTGGACACTTAAACAGGTCCTCAAGC	840
QY	942	AGAACTATTTCTATGCACTCAAGTTTACACAGGCTCAGCTGGCTTCAATATGNAATCT	1001
DB	841	AGAACTATTTCTATGCACTCAAGTTTACACAGGCTCAGCTGGCTTCAATATGNAATCT	900
QY	1002	TTCTGACATTTGATCAAGATGGAATACTTACAGCAGAGGAATTTATCCTGGCAATGCACCT	1061
DB	901	TTCTGACATTTGATCAAGATGGAATACTTACAGCAGAGGAATTTATCCTGGCAATGCACCT	960
QY	1062	CATTGATGTAGCTATGTCTGGCCAACTCACTGGCACCTGTCTCTGCCTCCAGAATACATTC	1121
DB	961	CATTGATGTAGCTATGTCTGGCCAACTCACTGGCACCTGTCTCTGCCTCCAGAATACATTC	1020
QY	1122	ACCTTCTTTTGAAGAGTTGATCTGGCAGTGGTATATCTGTCTCATAGCTCAACATCTGT	1181
DB	1021	ACCTTCTTTTGAAGAGTTGATCTGGCAGTGGTATATCTGTCTCATAGCTCAACATCTGT	1080
QY	1182	AGATCAGAGGCTACACAGAGGAACCACTTTTGAAGATGAACAACAACAAATTAGAAGAA	1241
DB	1081	AGATCAGAGGCTACACAGAGGAACCACTTTTGAAGATGAACAACAACAATTAGAAGAA	1140
QY	1242	ATTACCTGTAACTTTGAAGATAAGAACGGGAGAACTTTGAACGTGGCAACTGGAAC	1301
DB	1141	ATTACCTGTAACTTTGAAGATAAGAACGGGAGAACTTTGAACGTGGCAACTGGAAC	1200
QY	1302	GGAGAAAACGAAGCAAGCTCTCTGGAAACAGCAGCGCAAGGACGAGGCGCTGGCCCA	1361
DB	1201	GGAGAAAACGAAGCAAGCTCTCTGGAAACAGCAGCGCAAGGACGAGGCGCTGGCCCA	1260
QY	1362	GCTGGAGCGGGCGGAGCAGAGGAAGGAGCGTGGAGCCAGGAGCAGAGCGCAAGAG	1421
DB	1261	GCTGGAGCGGGCGGAGCAGAGGAAGGAGCGTGGAGCCAGGAGCAGAGCGCAAGAG	1320
QY	1422	ACAACTGGAACTGGGAAAGCAACTGGAAAAGCAGCGGGAGCTAGAAACGAGAGAGGA	1481
DB	1321	ACAACTGGAACTGGGAAAGCAACTGGAAAAGCAGCGGGAGCTAGAAACGAGAGAGGA	1380
QY	1482	GGAGAGGAGGAAAAGAAATTTGAGAGCGCAGAGCGTGCAAAACGGGAACTTTGAAAGGCAACG	1541
DB	1381	GGAGAGGAGGAAAAGAAATTTGAGAGCGCAGAGCGTGCAAAACGGGAACTTTGAAAGGCAACG	1440
QY	1542	ACAACTTGAGTGGGACGGAATTCGAAGGCAAGAACTTACTAAATCAAGAAAAACAAAGAA	1601
DB	1441	ACAACTTGAGTGGGACGGAATTCGAAGGCAAGAACTTACTAAATCAAGAAAAACAAAGAA	1500
QY	1602	AGAGGACATAGTTGTACTGAAAAGCAAGAAAAGACTTTTGGAAATTTGAAATTAGAAGCTCT	1661
DB	1501	AGAGGACATAGTTGTACTGAAAAGCAAGAAAAGACTTTTGGAAATTTGAAATTAGAAGCTCT	1560
QY	1662	AAATGATAAAAAGCATCAACTAGAAGGGGAAACTTCAAGATATCAGATGTGATTCAGCCAC	1721

DB	1561	AAATGATAAAAAGCATCAACTAGAAGGGAAACTTCAAGATATCAGATGTGATTCAGCCAC	1620
QY	1722	CCAAAGGCAAGAAATTTGAGAGCACAAAACAAATCTAGAGAGTTTGAGAAATTCGCGAAATCAC	1781
DB	1621	CCAAAGGCAAGAAATTTGAGAGCACAAAACAAATCTAGAGAGTTTGAGAAATTCGCGAAATCAC	1680
QY	1782	CCATCTACAGCAACAAATTTACAGGAATCTCAGCAAAATGCTTGGAGAGCTTTATTCAGAAAA	1841
DB	1681	CCATCTACAGCAACAAATTTACAGGAATCTCAGCAAAATGCTTGGAGAGCTTTATTCAGAAAA	1740
QY	1842	ACAGATATCTCAATGACCAATTTAAACACAGTTTCAGCAGAACAGTTTGCACAGAGATTCCT	1901
DB	1741	ACAGATATCTCAATGACCAATTTAAACACAGTTTCAGCAGAACAGTTTGCACAGAGATTCCT	1800
QY	1902	TGTTACACTTTAAAAGAGCGCTTTAGAAGCAAAAAGAACTAGCTCGGCAGCACCTACAGAGCCA	1961
DB	1801	TGTTACACTTTAAAAGAGCGCTTTAGAAGCAAAAAGAACTAGCTCGGCAGCACCTACAGAGCCA	1860
QY	1962	ACTGATGAAGTGGAGAAAAGAACTAGATCAAAAACCTACAGAGATTTGATATTTCAATAA	2021
DB	1861	ACTGATGAAGTGGAGAAAAGAACTAGATCAAAAACCTACAGAGATTTGATATTTTCAATAA	1920
QY	2022	TCAGCTGAAGGAACTTAAGAGAAATACACAATAAGCAACAACCTCCAGAAGCAAAAGTCCAT	2081
DB	1921	TCAGCTGAAGGAACTTAAGAGAAATACACAATAAGCAACAACCTCCAGAAGCAAAAGTCCAT	1980
QY	2082	GGAGCTGAAACGACTGAAAACAGAAAAGAAACAAAGAACGAAAGATCATAGAAATTTAGAAAA	2141
DB	1981	GGAGCTGAAACGACTGAAAACAGAAAAGAAACAAAGAACGAAAGATCATAGAAATTTAGAAAA	2040
QY	2142	AAAAGAAAGCCCAAAGACGAGCTCAGGAAGGGGAACAGCAGTGGCTGGAGCATGTGCA	2201
DB	2041	AAAAGAAAGCCCAAAGACGAGCTCAGGAAGGGGAACAGCAGTGGCTGGAGCATGTGCA	2100
QY	2202	CGAGGACGAGCATCAGAGCAACCAAGAAAACCTCCAGAAAGGGAACCTGAAAAGGGA	2261
DB	2101	CGAGGAGCAGCATCAGAGCAACCAAGAAAACCTCCAGAAAGGGAACCTGAAAAGGGA	2160
QY	2262	GGAGAGTGTCAAAAAGAGGATGGCGAGAAAAAGGCAAAACAGGAAGCACAAGACAAGCT	2321
DB	2161	GGAGAGTGTCAAAAAGAGGATGGCGAGAAAAAGGCAAAACAGGAAGCACAAGACAAGCT	2220
QY	2322	GGGTGGCTTTTTCATCAACCAAGAACCAAGCTTAAGCAGCTGTCCAGGCAACCTGGTC	2381
DB	2221	GGGTGGCTTTTTCATCAACCAAGAACCAAGCTTAAGCAGCTGTCCAGGCAACCTGGTC	2280
QY	2382	CACCTCGAAAAAAGTCCACTTTACCATTCTGCAACGAAAAATGTAAAAGTGGTGATTA	2441
DB	2281	CACCTCGAAAAAAGTCCACTTTACCATTCTGCAACGAAAAATGTAAAAGTGGTGATTA	2340
QY	2442	CCGGGCACTGTACCCCTTTTGAATCCAGAAGCCATGATGAAATCACTATCCAGCCAGGAGA	2501
DB	2341	CCGGGCACTGTGTACCCCTTTTGAATCCAGAAGCCATGATGAAATCACTATCCAGCCAGGAGA	2400
QY	2502	CATAGTCAAT-----GGTGGATGAAAAGCCAAAACCTGGAGAACCCGGCTGGCT	2546
DB	2401	CATAGTCAATGTTTAAAGGGGAATGGGTGGATGAAAAGCCAAAACCTGGAGAACCCGGCTGGCT	2460
QY	2547	TGGAGGAAATTTAAAGGAAAGACAGGCTGGTCCCTGCAAACTATGCAAGAGAAATCCC	2606
DB	2461	TGGAGGAAATTTAAAGGAAAGACAGGCTGGTCCCTGCAAACTATGCAAGAGAAATCCC	2520
QY	2607	AGAAAATGAGGTTCCCGCTCCAGTGAAACCAAGTGAATTAATCAACATCTGCCCTGCC	2666
DB	2521	AGAAAATGAGGTTCCCGCTCCAGTGAAACCAAGTGAATTAATCAACATCTGCCCTGCC	2580
QY	2667	CAAACTGGCTTGGTGAGACCCCGCCCTTTTGGCAGTAACCTTTCAGAGCCCTCCAC	2726
DB	2581	CAAACTGGCTTGGTGAGACCCCGCCCTTTTGGCAGTAACCTTTCAGAGCCCTCCAC	2640
QY	2727	GACCCCTAATAACTGGGCGGACTTCAGCTCCACCTGGGCCACACAGCAAGATGAGAAACC	2786

Db 2641 GACCCCTAATACTGGCGCAGCTTCAGCTCCAGTGGCCACCAGCAGCAATGAGAAACC 2700
Qy
2787 AGAAACGGATAAAGTGGATGCAATGGGACGCCAGCCCTCTCTCAACGGTTCGAAGTCCCG 2846
Db 2701 AGAAACGGATAAAGTGGATGCAATGGGACGCCAGCCCTCTCTCAACGGTTCGAAGTCCCG 2760
Qy 2847 CCAGTTAAGGCAGAGTCCGCTTTACTCCAGCCAGCGCACTGGCTCTCCCGCTCTCC 2906
Db 2761 CCAGTTAAGGCAGAGTCCGCTTTACTCCAGCCAGCGCACTGGCTCTCTCCCGCTCTCC 2820
Qy 2907 TGTGCTAGGCCAGAGTGAAGAGTGGAGGGCTTACAAGCTCAAGCCCTATATCTTTGGAG 2966
Db 2821 TGTGCTAGGCCAGAGTGAAGAGTGGAGGGCTTACAAGCTCAAGCCCTATATCTTTGGAG 2880
Qy 2967 AGCAAAAAAGCAACCACTTAATTTTAAACAAAATGATGTCATCACGTCCTGGAACA 3026
Db 2881 AGCAAAAAAGCAACCACTTAATTTTAAACAAAATGATGTCATCACGTCCTGGAACA 2940
Qy 3027 GCAAGACATGTGTGTGTGGAGAGTTCAGGTCAAGAGGGTGTGGTCCCAAGTCTTA 3086
Db 2941 GCAAGACATGTGTGTGTGGAGAGTTCAGGTCAAGAGGGTGTGGTCCCAAGTCTTA 3000
Qy 3087 CGTGAACCTCAATTCAGGGCCCAATAAGGAGTCTACAAGCATGGAATCTGGTCTTTCAGA 3146
Db 3001 CGTGAACCTCAATTCAGGGCCCAATAAGGAGTCTACAAGCATGGAATCTGGTCTTTCAGA 3060
Qy 3147 GAGTCTGTAGTCTAAAGGAGTAGCTCTCAGCAGCAAGCCGGTGTGTTCCGGGAGA 3206
Db 3061 GAGTCTGTAGTCTAAAGCAGTAGCTCTCAGCAGCAAGCCGGTGTGTTCCGGGAGA 3120
Qy 3207 A----- 3207
Db 3121 AGAATTTATGTCATGTACACTTACGAGAGTTCTGAGCAAGGAGATTTAACCTTTTCAGCA 3180
Qy 3208 ----- 3207
Db 3181 AGGGGATGTATTTGGTTACCAAGAAAGATGTGTGACTGTGTGGACAGGAACAGTGGCGGA 3240
Qy 3208 ----- 3207
Db 3241 CAAGGCGGAGTCTTCCCTTCTAATATGTGAGGCTTAAAGATTTCAGAGGGCTCTGGAAC 3300
Qy 3208 ----- GAAATTTCCCGAGTTATTCCTCATTA 3233
Db 3301 TGTGCGAAACAGGAGATTAGAAACAACTGAAATTTGCCAGGTTATGCTCATTA 3360
Qy 3234 CACCGCCACCGCCCGGAGCAGCTCACTCTCGCCCTCTGGTCAAGTATTTGATCCGAAA 3293
Db 3361 CACCGCCACCGCCCGGAGCAGCTCACTCTCGCCCTCTGGTCAAGTATTTGATCCGAAA 3420
Qy 3294 AAGAAGCCAGGTGGATGGTGGGAAAGAGAGTGCAGGACAGTGGGGAAGAGCCCGAGAT 3353
Db 3421 AAGAAGCCAGGTGGATGGTGGGAAAGAGAGTGCAGGACAGTGGGGAAGAGCCCGAGAT 3480
Qy 3354 AGCTGGTTCCTCAGCTAATTTATGATGAGTCTTAAGCCCTGGGACGAGCAAAATCACTCC 3413
Db 3481 AGCTGGTTCCTCAGCTAATTTATGATGAGTCTTAAGCCCTGGGACGAGCAAAATCACTCC 3540
Qy 3414 AACAGAGCCACCTAAGTCAACAGCATTTAGCGGAGTGTGCGCAGGTGATTTGGGATGTACGA 3473
Db 3541 AACAGAGCCACCTAAGTCAACAGCATTTAGCGGAGTGTGCGCAGGTGATTTGGGATGTACGA 3600
Qy 3474 CTAACCGCGCAGATGAGTGGCTTTCAACAGGGCCAGATCATCAACGTCTCT 3533
Db 3601 CTAACCGCGCAGATGAGTGGCTTTCAACAGGGCCAGATCATCAACGTCTCT 3660
Qy 3534 CAACAAGGAGCCCTCAGCTGGTGGAAAGGAGAGTCAATGGCAAGTGGGGCTCTTCCC 3593
Db 3661 CAACAAGGAGCCCTCAGCTGGTGGAAAGGAGAGTCAATGGCAAGTGGGGCTCTTCCC 3720
Qy 3594 ATCCAAATTTATGAGTGTGACCAAGACATGGACCCCAAGCCAGCAATG 3641
Db 3721 ATCCAAATTTATGAGTGTGACCAAGACATGGACCCCAAGCCAGCAATG 3768

RESULT 3

US-10-287-436A-685
; Sequence 685, Application US/10287436A
; Publication No. US20050202421A1
; GENERAL INFORMATION:
; APPLICANT: CHILDREN'S HOSPITAL MEDICAL CENTER
; TITLE OF INVENTION: METHOD FOR DIAGNOSIS AND TREATMENT OF
; TITLE OF INVENTION: RHEUMATOID ARTHRITIS
; FILE REFERENCE: 10872.514696
; CURRENT APPLICATION NUMBER: US/10/287,436A
; CURRENT FILING DATE: 2002-10-31
; PRIOR APPLICATION NUMBER: US 60/336,220
; PRIOR FILING DATE: 2001-10-31
; NUMBER OF SEQ ID NOS: 1446
; SOFTWARE: FASTSEQ for Windows Version 4.0
; SEQ ID NO 685
; LENGTH: 7247
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-287-436A-685

Query Match 63.2%; Score 3285.6; DB 9; Length 7247;

Best Local Similarity 93.8%; Pred. No. 0;

Matches 3536; Conservative 0; Mismatches 4; Indels 228; Gaps 2;

Qy 102 GCGTCCCTCCAGCGCGGTGAGCGGCACTGATTTGTCCCTGGGGCGGCGAGCGGAC 161
Db 1 GCGTCCCTCCAGCGCGGTGAGCGGCACTGATTTGTCCCTGGGGCGGCGAGCGGAC 60
Qy 162 CGCGCCGAGATGAGGCGTCCGATTTAGCAAGGTAAAGTAACAGAACCATGCTCAGTTTCC 221
Db 61 CGCGCCGAGATGAGGCGTCCGATTTAGCAAGGTAAAGTAACAGAACCATGCTCAGTTTCC 120
Qy 222 AACACCTTTTGTGGCAGCCTTGATATCTGGGCCATAAAGTGTAGAGAAAGAGCGAGCA 281
Db 121 AACACCTTTTGTGGCAGCCTTGATATCTGGGCCATAAAGTGTAGAGAAAGAGCGAGCA 180
Qy 282 TGNATGAGCTATCCAGTATTTAAAGCAATATCTGGATTCATTTCTGGTGTATCAAGCTAG 341
Db 181 TGNATGAGCTATCCAGTATTTAAAGCAATATCTGGATTCATTTCTGGTGTATCAAGCTAG 240
Qy 342 AAAATTTTTTTTCAATCTGGGTACTCTCAACCTGTTTGTAGCAGATATGGGCACTAGC 401
Db 241 AAAATTTTTTTTCAATCTGGGTACTCTCAACCTGTTTGTAGCAGATATGGGCACTAGC 300
Qy 402 TGACATGAATTAATGATGGAGAAATGGATCAAGTGGAGTTTTTCCATAGCTATGAACCTAT 461
Db 301 TGACATGAATTAATGATGGAGAAATGGATCAAGTGGAGTTTTTCCATAGCTATGAACCTAT 360
Qy 462 CAAACTGAAGCTACAGGATATCAGCTACCTCTGCACTTCCCTCTGATGATGAACAGCA 521
Db 361 CAAACTGAAGCTACAGGATATCAGCTACCTCTGCACTTCCCTCTGATGATGAACAGCA 420
Qy 522 ACCAGTTGCTATTTCTAGCGCACAGCATTTGGTATGGGAGGTATGCCAGCATGCCACC 581
Db 421 ACCAGTTGCTATTTCTAGCGCACCAATTTGGTATGGGAGGTATGCCAGCATGCCACC 480
Qy 582 GCTTACAGCTGTTTGTCTCCAGTGCCCAATGGGATCCATTTCCAGTTTGTGGATGTCTCAAC 641
Db 481 GCTTACAGCTGTTTGTCTCCAGTGCCCAATGGGATCCATTTCCAGTTTGTGGATGTCTCAAC 540
Qy 642 CCTAGTATCTTCTGTTCCCAACAGCATGTGCCCCCTCTGCTAAACGGGGCTCCCTCTGT 701
Db 541 CCTAGTATCTTCTGTTCCCAACAGCATGTGCCCCCTCTGCTAAACGGGGCTCCCTCTGT 600
Qy 702 TATACAACTCTGCTGCAATTTGCTCATCTCGAGCCACATTTGCCAAGAGTTCTTCTT 761
Db 601 TATACAACTCTGCTGCAATTTGCTCATCTCGAGCCACATTTGCCAAGAGTTCTTCTT 660
Qy 762 TAGTAGATCTGGTCCAGGGTCAAACTAAATTAACAAAGGCACAGTCATTGTA 821

Dd	661	TAGTAGATCTGCTCCAGGGTCAAACTAAACATAAAATTAACAAAAGGCACAGTCAATTGA	720
Qy	822	TGTGGCCAGTGTCCACCAAGTGGCAGAGTGGGCTGTTCTCTCAGTCATCAAGACTGAAATA	881
Dd	721	TGTGGCCAGTGTCCACCAAGTGGCAGAGTGGGCTGTTCTCTCAGTCATCAAGGCTGAATA	780
Qy	882	CAGGCAATTATTCAATAGTCATGACAAACTATAGTGGAGACACTTAAACAGTCCCAAGC	941
Dd	781	CAGGCAATTATTCAATAGTCATGACAAACTATAGTGGAGACACTTAAACAGTCCCAAGC	840
Qy	942	RAGAACTATTCTTATGCAAGTCAAGTTTACCACAGGCTCAGCTGGCTTCAATATGAATCT	1001
Dd	841	AAAGAACTATTCTTATGCAAGTCAAGTTTACCACAGGCTCAGCTGGCTTCAATATGAATCT	900
Qy	1002	TTCTGACATTTGATCAAGATGGAACCTTACAGCAGAGGAATTTATCTGGCAATGCACCT	1061
Dd	901	TTCTGACATTTGATCAAGATGGAACCTTACAGCAGAGGAATTTATCTGGCAATGCACCT	960
Qy	1062	CATTGATGTAGCTATGCTGGCCCAACCACTGCCACTGCTCTGCTCCAGAAATCATTTCC	1121
Dd	961	CATTGATGTAGCTATGCTGGCCCAACCACTGCCACTGCTCTGCTCCAGAAATCATTTCC	1020
Qy	1122	ACCTTCTTTTAGAAGAGTTTCGATCTGGCAGTGGTATATCTGTATAAGCTCAACATCTGT	1181
Dd	1021	ACCTTCTTTTAGAAGAGTTTCGATCTGGCAGTGGTATATCTGTATAAGCTCAACATCTGT	1080
Qy	1182	AGATCAGAGGCTACCAAGGAAACCAAGTTTATAGAAGTGAACCAACAAATTAGAAAGAA	1241
Dd	1081	AGATCAGAGGCTACCAAGGAAACCAAGTTTATAGAAGTGAACCAACAAATTAGAAAGAA	1140
Qy	1242	ATTACTGTAAAGTTTGAAGATAAGAGCGGGAGAACTTTGAACGTGGCAACTGGAACCT	1301
Dd	1141	ATTACTGTAAAGTTTGAAGATAAGAGCGGGAGAACTTTGAACGTGGCAACTGGAACCT	1200
Qy	1302	GGAGAAACGAAGCAAGCTCTCTTGAACAGAGCGCAAGGAGCAGAGCGCTGGCCCA	1361
Dd	1201	GGAGAAACGAAGCAAGCTCTCTTGAACAGAGCGCAAGGAGCAGAGCGCTGGCCCA	1260
Qy	1362	GCTGAGCGGGCGGAGCAGAGAGGAGGAGCGGTGAGCGCCAGGAGCAAGAGCGCAAAAG	1421
Dd	1261	GCTGAGCGGGCGGAGCAGAGAGGAGGAGCGGTGAGCGCCAGGAGCAAGAGCGCAAAAG	1320
Qy	1422	ACAACTGGAACTGGAGAACCACTGGAAGAGCAGCGGGAGCTTAGAACCGCAGAGAGGA	1481
Dd	1321	ACAACTGGAACTGGAGAACCACTGGAAGAGCAGCGGGAGCTTAGAACCGCAGAGAGGA	1380
Qy	1482	GGAGAGGAGAAAGAAATTTGAGAGCGAGAGGCTGCAAAACGGAACTTTGAAAGGCAAG	1541
Dd	1381	GGAGAGGAGAAAGAAATTTGAGAGCGAGAGGCTGCAAAACGGAACTTTGAAAGGCAAG	1440
Qy	1542	ACAACTGGAGTGGAAACGGAATCGAAGGCAAGAACTTACTAAATCAAAGAAACAAAGAAC	1601
Dd	1441	ACAACTGGAGTGGAAACGGAATCGAAGGCAAGAACTTACTAAATCAAAGAAACAAAGAAC	1500
Qy	1602	AGAGACATAGTTTACTGAAAGCAAGAAAGAACTTTGGAAATTTGAAATTTAGAGCTCT	1661
Dd	1501	AGAGACATAGTTTACTGAAAGCAAGAAAGAACTTTGGAAATTTGAAATTTAGAGCTCT	1560
Qy	1662	AAATGATAAAGATCAACTAGAGGGAACTTCAAGATATCAGATGTGCAATTTGACAC	1721
Dd	1561	AAATGATAAAGATCAACTAGAGGGAACTTCAAGATATCAGATGTGCAATTTGACAC	1620
Qy	1722	CCAAAGGCAAGAAATTTAGAGCAGCAACAAATCTAGAGAGTTGAGAAATTTGCCGAAATCAC	1781
Dd	1621	CCAAAGGCAAGAAATTTAGAGCAGCAACAAATCTAGAGAGTTGAGAAATTTGCCGAAATCAC	1680
Qy	1782	CCATCTACAGCAACAAATTTACAGGAATCTCAGCAATAGCTTGGAGACTTATTCCAGAAA	1841
Dd	1681	CCATCTACAGCAACAAATTTACAGGAATCTCAGCAATAGCTTGGAGAGCTTATTCCAGAAA	1740
Qy	1842	ACAGATACTCAATGACCAATTAACCAAGTTTACAGAGAAACAGTTTCCACAGAGATTCACT	1901
Dd	1741	ACAGATACTCAATGACCAATTAACCAAGTTTACAGAGAAACAGTTTCCACAGAGATTCACT	1800
Qy	1902	TGTTACACTTAAAAAGAGCCTTTAGAAAGCAAAAGAACTAGCTCGGAGGACCTTACGAGACCA	1961
Dd	1801	TGTTACACTTAAAAAGAGCCTTTAGAAAGCAAAAGAACTAGCTCGGAGGACCTTACGAGACCA	1860
Qy	1962	ACTGATCAAGTGGAGAAAGAACTTAGATCAAAACTACAGGAGATTGATATTTTCAATAA	2021
Dd	1861	ACTGATCAAGTGGAGAAAGAACTTAGATCAAAACTACAGGAGATTGATATTTTCAATAA	1920
Qy	2022	TCAGCTGAAGAACTTAAGAGAAATACACAAATAAGCAACAACCTCCAGAGCAAAAAGTCCAT	2081
Dd	1921	TCAGCTGAAGAACTTAAGAGAAATACACAAATAAGCAACAACCTCCAGAGCAAAAAGTCCAT	1980
Qy	2082	GGAGCTGAACGACTGAAGAAACAGAAAGAAACAAAGATCATAGAAATTTAGAAAAACA	2141
Dd	1981	GGAGCTGAACGACTGAAGAAACAGAAAGAAACAAAGATCATAGAAATTTAGAAAAACA	2040
Qy	2142	AAAAGAAAGCCCAAGACGAGCTCAGAAAGGACAAAGCAGTGGCTGGAGCATGTGCA	2201
Dd	2041	AAAAGAAAGCCCAAGACGAGCTCAGAAAGGACAAAGCAGTGGCTGGAGCATGTGCA	2100
Qy	2202	GCAGAGGACGAGCATCAGAGACCAAGAAAACTCCAAGAGGAGAAAACTGAAAAAGGA	2261
Dd	2101	GCAGAGGACGAGCATCAGAGACCAAGAAAACTCCAAGAGGAGAAAACTGAAAAAGGA	2160
Qy	2262	GGAGAGTGTCAAAAAGAGGATGGCGAGGAAAAAGGCAAAACAGGAGCACAAGCAAGCT	2321
Dd	2161	GGAGAGTGTCAAAAAGAGGATGGCGAGGAAAAAGGCAAAACAGGAGCACAAGCAAGCT	2220
Qy	2322	GGTTCGGCTTTTCCATCAACCAAGAACCAAGCAGCTGTCAGGAGCAACCTGGTC	2381
Dd	2221	GGTTCGGCTTTTCCATCAACCAAGAACCAAGCAGCTGTCAGGAGCAACCTGGTC	2280
Qy	2382	CACTCAGAGAAAGGTCCACTTTTACCATTTCTGCAACAGGAAAAATGTAAGTGGTGTATTA	2441
Dd	2281	CACTCAGAGAAAGGTCCACTTTTACCATTTCTGCAACAGGAAAAATGTAAGTGGTGTATTA	2340
Qy	2442	CGGGCACTGTACCCCTTTGAAATCCAGAACCATGATGAAATCACTATCCAGCCAGGAGA	2501
Dd	2341	CGGGCACTGTACCCCTTTGAAATCCAGAACCATGATGAAATCACTATCCAGCCAGGAGA	2400
Qy	2502	CATAGTCAT-----GGTGGATGAAAGCACAACCTGGAGAAACCGCGCTGGCT	2546
Dd	2401	CATAGTCATGGTTTAAAGGGGAATGGGTGGATGAAAGCACAACCTGGAGAAACCGCGCTGGCT	2460
Qy	2547	TGGAGGAGAAATTAAGAGGAAAGACAGGGTGGTTCCTGCAAACTATGAGAGAAAAATCCC	2606
Dd	2461	TGGAGGAGAAATTAAGAGGAAAGACAGGGTGGTTCCTGCAAACTATGAGAGAAAAATCCC	2520
Qy	2607	AGAAAATGAGGTTCCCGCTCCAGTGAACCAAGTGAATGATTCACATCTGCCCTGGCCC	2666
Dd	2521	AGAAAATGAGGTTCCCGCTCCAGTGAACCAAGTGAATGATTCACATCTGCCCTGGCCC	2580
Qy	2667	CAAACTGGCTTGGTGAAGACCCCGCTTTTGGCAGTAACTCTTCCAGAGCCCTCCAC	2726
Dd	2581	CAAACTGGCTTGGTGAAGACCCCGCTTTTGGCAGTAACTCTTCCAGAGCCCTCCAC	2640
Qy	2727	GAACCTTAATACTGGGCGCACTTCAAGTCAAGTGGGCGCCACAGCAAGAAATGAGAAACC	2786
Dd	2641	GAACCTTAATACTGGGCGCACTTCAAGTCAAGTGGGCGCCACAGCAAGAAATGAGAAACC	2700
Qy	2787	AGAAAACGATTAACCTGGATGATGGGAGCCCGCTCTCTCACCGTTCCAGGTGGCGG	2846
Dd	2701	AGAAAACGATTAACCTGGATGATGGGAGCCCGCTCTCTCACCGTTCCAGGTGGCGG	2760
Qy	2847	CCAGTTAAGGCAAGGTCGCTTTTACTCCAGCCACGGCCACTGGCTCTCTCCCGCTCTCC	2906
Dd	2761	CCAGTTAAGGCAAGGTCGCTTTTACTCCAGCCACGGCCACTGGCTCTCTCCCGCTCTCC	2820
Qy	2907	TGTGTAGGCCAGGGTGAAAAAGGTGGAGGGGTACAAGCTCAAGCCCTATATCTTCTTGGAG	2966
Dd	2821	TGTGTAGGCCAGGGTGAAAAAGGTGGAGGGGTACAAGCTCAAGCCCTATATCTTCTTGGAG	2880


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QY 2967 AGCCAAAAGAACCACTTAATTTTAAACAAAATGATGTCATCACCGTCTCTGGAACA 3026
Db 2881 AGCCAAAAGAACCACTTAATTTTAAACAAAATGATGTCATCACCGTCTCTGGAACA 2940
QY 3027 GCAAGACATGTGTGTGGTGGAGAAAGTTCAAGGTGAGAGGGTGGTTCCTCCCAAGTCTTA 3086
Db 2941 GCNAGACATGTGTGTGGTGGAGAAAGTTCAAGGTGAGAGGGTGGTTCCTCCCAAGTCTTA 3000
QY 3087 CGTGAACCTCAATTCAGGGCCCAATAAGAAAGTCTCAAGCATGATCTGGTCTTCAGA 3146
Db 3001 CGTGAACCTCAATTCAGGGCCCAATAAGAAAGTCTCAAGCATGATCTGGTCTTCAGA 3060
QY 3147 GAGTCTGTAGTCTAAAGGAGTCTCTCAGCAGCAACCGGTCTGTTTCGGGAGA 3206
Db 3061 GAGTCTGTAGTCTAAAGGAGTCTCTCAGCAGCAACCGGTCTGTTTCGGGAGA 3120
QY 3207 A----- 3207
Db 3121 AGAATTTATGCCATGTACACTTACGAGAGTCTCTGAGCAAGGAGATTTAACTTTTCAGCA 3180
QY 3208 ----- 3207
Db 3181 AGGGATGTGATTTTGGTTTACCAAGAAAGATGGTGAAGTCTGAGTGGACAGAAACAGTGGGCGA 3240
QY 3208 ----- 3207
Db 3241 CAAGCGCGGAGTCTTCCCTTCTAACTATGTAGAGCTTTAAAGATTCAAGAGGCTCTGGAAC 3300
QY 3208 -----GAAATTTGCCCAGGTATTTGCCTCAT 3233
Db 3301 TGTGGGAAACAGGGAGTTTAGGAAATAAACCTGAAATTTGCCAGGTATTTGCTTCATA 3360
QY 3234 CACCGCCACCGGCCCGAGCAGTCACTCTCGCCCTCTGGTCAAGTGAATTTGATCCGAAA 3293
Db 3361 CACCGCCACCGGCCCGAGCAGTCACTCTCGCCCTCTGGTCAAGTGAATTTGATCCGAAA 3420
QY 3294 AAAGAACCCAGGTGGATGGTGGNAGGAGAGCTGCAAGCAGCTGGGAAAGCGCCAGAT 3353
Db 3421 AAAGAACCCAGGTGGATGGTGGNAGGAGAGCTGCAAGCAGCTGGGAAAGCGCCAGAT 3480
QY 3354 AGGCTGGTCTCCAGCTAATTTATGTAAAGCTTTCTAAGCCCTGGGACGAGCAAAATCACTCC 3413
Db 3481 AGGCTGGTCTCCAGCTAATTTATGTAAAGCTTTCTAAGCCCTGGGACGAGCAAAATCACTCC 3540
QY 3414 AACAGACCCACTAAGTCAACAGCATTTAGCGGCAGTGTGCCAGGTGATTTGGGATGTACGA 3473
Db 3541 AACAGACCCACTAAGTCAACAGCATTTAGCGGCAGTGTGCCAGGTGATTTGGGATGTACGA 3600
QY 3474 CTACACCGCGAGATGACGATGAGCTGGGCTTCAACNAGGGCCAGATCATCAACGTCCT 3533
Db 3601 CTACACCGCGAGATGACGATGAGCTGGGCTTCAACNAGGGCCAGATCATCAACGTCCT 3660
QY 3534 CAACAAGGAGGACCTCACTGCTGGAAAGGAGAAAGTCAATGGACAAGTGGGGCTCTTTCCC 3593
Db 3661 CAACAAGGAGGACCTCACTGCTGGAAAGGAGAAAGTCAATGGACAAGTGGGGCTCTTTCCC 3720
QY 3594 ATCCAATTTATGTGAAGTGTACCAAGATGAGTGGACCCAGCCAGCAATG 3641
Db 3721 ATCCAATTTATGTGAAGTGTACCAAGATGAGTGGACCCAGCCAGCAATG 3768
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RESULT 4
US-10-450-763-20567
; Sequence 20567, Application US/10450763
; Publication No. US2005019675441
; GENERAL INFORMATION:
; APPLICANT: Hyseq, Inc
; TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES
; FILE REFERENCE: 790CIE3/US
; CURRENT APPLICATION NUMBER: US/10/450,763
; CURRENT FILING DATE: 2003-06-11
; PRIOR APPLICATION NUMBER: PCT/US01/08631
; PRIOR FILING DATE: 2001-03-30
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; PRIOR APPLICATION NUMBER: 09/540,217
; PRIOR FILING DATE: 2000-03-31
; PRIOR APPLICATION NUMBER: 09/649,167
; PRIOR FILING DATE: 2000-08-23
; NUMBER OF SEQ ID NOS: 60736
; SOFTWARE: Custom
; SEQ ID NO 20567
; LENGTH: 7435
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SIMILAR
; LOCATION: (263)...(2356)
; OTHER INFORMATION: 66% homologous to Rattus norvegicus EH-domain/SH3-domain
; OTHER INFORMATION: containing protein, accession number AF132672, Smith-Waterman Score
; OTHER INFORMATION: 2106.
US-10-450-763-20567

Query Match 59.4%; Score 3090.6; DB 9; Length 7435;
Best Local Similarity 91.9%; Pred. No. 0;
Matches 3573; Conservative 0; Mismatches 54; Indels 260; Gaps 19;

QY 15 GTACGGCGGCTCCGAGGAAGATCCGAGCGGGCTCCGGAGCGGACAGAGAGCGGGCG 74
Db 70 GTACGGCGGCTCCGAGGAAGATCCGAGCGGGCTCCGGAGCGGACAGAGAGCGGGCG 129
QY 75 GGGATGTTGTCGGGGCTCGCGGTCTCTGCGTCCCTCCAGCGGCGCTGAGCGGCACTGA 134
Db 130 GGGATGTTGTCGGGGCTCGCGGTCTCTGCGTCCCTCCAGCGGCGCTGAGCGGCACTGA 189
QY 135 TTTTGTCCCTGGGGCGGCGAGCGCGACCCGCCCGGAGATGAGGGCTCGATTAGCAAGTAA 194
Db 190 TTTTGTCCCTGGGGCGGCGAGCGCGACCCGCCCGGAGATGAGGGCTCGATTAGCAAGTAA 249
QY 195 AAGTAACAGAAACATGGCTCAGTTTCCAAACACTTTTGGTGGAGCGCTGGATATCTGGGC 254
Db 250 AAGTAACAGAAACATGGCTCAGTTTCCAAACACTTTTGGTGGAGCGCTGGATATCTGGGC 309
QY 255 CATTAACGTAGAGGAAGAGCGAAGCATGATCAGCAGTTCATAGTTTAAAGCCAAATATC 314
Db 310 CATTAACGTAGAGGAAGAGCGAAGCATGATCAGCAGTTCATAGTTTAAAGCCAAATATC 369
QY 315 TGGATTCATTACTTGGTGATCAAGCTAGAAACTTTTTCATCTGGGTTTACCTCAACC 374
Db 370 TGGATTCATTACTTGGTGATCAAGCTAGAAACTTTTTCATCTGGGTTTACCTCAACC 429
QY 375 TGTTTTAGCACAGATATGGGCACTAGCTGACATGAATAATGATGGAGAAATGGATCAAGT 434
Db 430 TGTTTTAGCACAGATATGGGCACTAGCTGACATGAATAATGATGGAGAAATGGATCAAGT 489
QY 435 GGAGTTTTCATAGCTATGAAACTTTATCAAACTGAAGCTCAAGGATATCAAGTACCCTC 494
Db 490 GGAGTTTTCATAGCTATGAAACTTTATCAAACTGAAGCTCAAGGATATCAAGTACCCTC 549
QY 495 TGCACTTCCCTCTGTCTATGAAACAGCAACCCAGTTTGTCTATTCTAGCGCACCAAGATTGG 554
Db 550 TGCACTTCCCTCTGTCTATGAAACAGCAACCCAGTTTGTCTATTCTAGCGCACCAAGATTGG 609
QY 555 TATGGGAGGTATCCGACAGATGCCCGCTTACAGCTGTGCTCCAGTGGCCCAATGGGATC 614
Db 610 TATGGGAGGTATCCGACAGATGCCCGCTTACAGCTGTGCTCCAGTGGCCCAATGGGATC 669
QY 615 CATT-CCAGTTTGGGAATGTCCTCAACCCCTAGTATCTTCTGTTCCCAAGCAGAGCTGTGC 673
Db 670 CATTCCAGTTTGGGAATGTCCTCAACCCCTAGTATCTTCTGTTCCCAAGCAGAGCTGTGC 729
QY 674 CCCCCCTGGCTAACGGGGCTCCCCCTGTATACAACTCTCTCCCTGCAATTTGCTCATCTG 733
Db 730 CCCCCCTGGCTAACGGGGCTCCCCCTGTATACAACTCTCTCCCTGCAATTTGCTCATCTG 789
QY 734 CAGCAACATGCCAAAGAGTTCTTCTTTTAGTAGATCTGGTCCAGGGTCAACCTAAACA 793
Db 790 CAGCAACATGCCAAAGAGTTCTTCTTTTAGTAGATCTGGTCCAGGGTCAACCTAAACA 849
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QY	794	CTAAATTACAAAGGCACAGTCAATTTGATGTGGCCAGTCTCCACC - AGTGCAGAGTGG	852	QY	1858	CAATTTAAAACAAAGTTTCAGCAGNA	CAGTTTTCACAGAGATTCACTTGTTCACACTTAAAG -	1916
DB	850	CTAAATTACAAAGGCACAGTCAATTTGATGTGGCCAGTCTCCACCAGTGGCAGAGTGG	909	DB	1930	CAATTTAAAACAAAGTTTCAGCAGNA	CAGTTTTCACAGAGATTCACTTGTTCACACTTAAAG -	1989
QY	853	GCTGTTCTCAGTCATCAAGACTGAATAC - - - GGCATTTATCAATAGTCATGACAA	909	QY	1917	AGCCTTTAGAAGCAAAAGAACTAGCTCGGCAGCACCTACGAGACCACTGGATGGAAGTGA	1976	
DB	910	GCTGTTCTCAGTCATCAAGACTGAATAC - - - GGCATTTATCAATAGTCATGACAA	969	DB	1990	AGCCTTTAGAAGCAAAAGAACTAGCTCGGCAGCACCTACGAGACCACTGGATGGAAGTGA	2049	
QY	910	AC - -TATGAGTGACACTT - -AACAGGTCCCAAGCAAGAACTATTCTTATGTCAGTCAAG	965	QY	1977	GAAGAAATCTAGATCAAAACTACAGGAGATTGATATTTTCAATATAGCTGAAGAACT	2036	
DB	970	ACTTAATGAGTGGACACTTTTAACAGGTCCCAAGCAAGAACTATTCTTATGTCAGTCAAG	1029	DB	2050	GAAGAAATCTAGATCAAAACTACAGGAGATTGATATTTTCAATATAGCTGAAGAACT	2109	
QY	966	TTTACCACAGGCTCAGCTGGCTTCAATATGGAATCTTTCTGACATTTGATCAAGATGGA	1025	QY	2037	RAGAGAAATACAAATAGCAAACTCCAGAGCAAAAGTCCATGGAGGCTGAAGACT	2096	
DB	1030	TTTACCACAGGCTCAGCTGGCTTCAATATGGAATCTTTCTGACATTTGATCAAGATGGA	1089	DB	2110	RAGAGAAATACAAATAGCAAACTCCAGAGCAAAAGTCCATGGAGGCTGAAGACT	2169	
QY	1026	ACTTACAGCAGAGGAATTTATCTTGGCAATGCACCTCATTGATGCTATGCTGGCCA	1085	QY	2097	GAACACAGAAAGAAACAAAGAACGAAAGATCATAGAAATTTAGAAAAACAAAAGAGAGGCCA	2156	
DB	1090	ACTTACAGCAGAGGAATTTATCTTGGCAATGCACCTCATTGATGCTATGCTGGCCA	1149	DB	2170	GAACACAGAAAGAAACAAAGAACGAAAGATCATAGAAATTTAGAAAAACAAAAGAGAGGCCA	2229	
QY	1086	ACCACCTGCCACCTGCTCCTCCAGAAATACATTTCCACCTTCTTTTAGAAGAGTTGATC	1145	QY	2157	AAAGCAGCTCAGGAAAGGGAACAAGCTGGCTGGAGCATGTGCAGCAGGAGGACGAGCA	2216	
DB	1150	ACCACCTGCCACCTGCTCCTCCAGAAATACATTTCCACCTTCTTTTAGAAGAGTTGATC	1209	DB	2230	AAAGCAGCTCAGGAAAGGGAACAAGCTGGCTGGAGCATGTGCAGCAGGAGGACGAGCA	2289	
QY	1146	TGGCAGTGTATATCTGTATAGCTCAACATCTGTAGATCAGAGGCTACGAGAGAAC	1205	QY	2217	TCAGAGACCNAGAAACTCCAGAGAGGAAACTGAAAAAGGAGGAGAGTGTCAAAA	2276	
DB	1210	TGGCAGTGTATATCTGTATAGCTCAACATCTGTAGATCAGAGGCTACGAGAGAAC	1269	DB	2290	TCAGAGACCNAGAAACTCCAGAGAGGAAACTGAAAAAGGAGGAGAGTGTCAAAA	2349	
QY	1206	AGTTTATAGAGTGAACAACTATTAGAAAAGAAATTTACCTGTAAACCTTTGAGATAA	1265	QY	2277	GAAGGATGSC - GAGGAAAAAGGCAACAGGAAGCAACAAGCAAGCTGGGTTCGGCTTTTCC	2335	
DB	1270	AGTTTATAGAGTGAACAACTATTAGAAAAGAAATTTACCTGTAAACCTTTGAGATAA	1329	DB	2350	GAAGGATGGCAGAGGAAAGGCAACAGGAAGCACAGCAAGCTGGGTTCGGCTTTTCC	2409	
QY	1266	GAAGCGGAGAACTTTGAACTGGCAACTGGAACTGGAGAAACGAAGGCAAGCTCTCCT	1325	QY	2336	ATCAACACAAAGAACAGCTAAGCTGTGCAGGACCTCTGGTCCACTGCAGAGAAAG	2395	
DB	1330	GAAGCGGAGAACTTTGAACTGGCAACTGGAACTGGAGAAACGAAGGCAAGCTCTCCT	1389	DB	2410	ATCAACACAAAGAACAGCTAAGCTGTGCAGGACCTCTGGTCCACTGCAGAGAAAG	2469	
QY	1326	GGAAACAGCGCAAGGACGAGAGCGCTTGGCCAGCTGGAGCGGCGGAGCAGAGAG	1385	QY	2396	GTCCACTTACCAATTTTCACAGGAAATGTAAAAAGTGTGTATTACCGGGCACTGTACC	2455	
DB	1390	GGAAACAGCGCAAGGACGAGAGCGCTTGGCCAGCTGGAGCGGCGGAGCAGAGAG	1449	DB	2470	GTCCACTTACCAATTTTCACAGGAAATGTAAAAAGTGTGTATTACCGGGCACTGTACC	2529	
QY	1386	GAAGAGCTGAGCGCCAGGACACAGAGCGCAAGACAACTGGAACCTGGAGAGCAACT	1445	QY	2456	CTTTTGAATCCAGAAAGCCATGATGAAATCACTATCCAGCAGGAGACATAGTCAAT - - - - -	2510	
DB	1450	GAAGAGCTGAGCGCCAGGACACAGAGCGCAAGACAACTGGAACCTGGAGAGCAACT	1509	DB	2530	CTTTTGAATCCAGAAAGCCATGATGAAATCACTATCCAGCAGGAGACATAGTCAATGTTA	2589	
QY	1446	GGAAACAGCGGAGCTAGAAACGGCAGAGAGGAGGAGAGGAGGAGGAGGAGGAGGAG	1505	QY	2511	----- -GGTGATGAAAGCCAAAACCTGGAGAACCCGGCTGGCTTGGAGAGAGATTAA	2560	
DB	1510	GGAAACAGCGGAGCTAGAAACGGCAGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG	1569	DB	2590	AAGCGGAATGGGTGGATGAAAGCCAACTGGAGAACCCGGCTGGCTTGGAGAGAGATTAA	2649	
QY	1506	GCAGAGGCTGCAAAACGGGAAC - -TGAAAGGCAACGACAACTTGTAGTGGGAACGGAATC	1564	QY	2561	RAGGAAAGACAGGTGGTTCCCTGCAAACTATGCAGAGAAAAATCCAGAAAAATGAGGTTT	2620	
DB	1570	GCAGAGGCTGCAAAACGGGAAC - -TGAAAGGCAACGACAACTTGTAGTGGGAACGGAATC	1629	DB	2650	RAGGAAAGACAGGTGGTTCCCTGCAAACTATGCAGAGAAAAATCCAGAAAAATGAGGTTT	2709	
QY	1565	GAAGCAAGAACTACTTAATCAAGAAACAAAGAACAGGAGCAGTGTGTACTGAAAG	1624	QY	2621	CCGCTCCAGTGAAACAGTGACTGATTCAAATCTGCCCTTCCCTCCCTCCCTCCCTCCCT	2680	
DB	1630	GAGAGTCCAACTACTTAATCAAGAAACAAAGAACAGGAGCAGTGTGTACTGAAAG	1689	DB	2710	CCGCTCCAGTGAAACAGTGACTGATTCAAATCTGCCCTTCCCTCCCTCCCTCCCTTGC	2769	
QY	1625	CAAGAAAAAGACTTTTGAATTTGAATTTAGAGCTCTAAATGATAAAAAAGCATCAACTAG	1684	QY	2681	GTGAGACCCCTCCCTCTTGGCAGTAACTCTTTCAGAGCCCTCCAGACCCCTTAATACT	2740	
DB	1690	CAAGAAAAAGACTTTTGAATTTGAATTTAGAGCTCTAAATGATAAAAAAGCATCAACTAG	1749	DB	2770	GTGAGACCCCTCCCTCTTGGCAGTAACTCTTTCAGAGCCCTCCAGACCCCTTAATACT	2829	
QY	1685	AGGGGAACTTCAAGATATCAGATGTCGATTTGACCAACCAAGGCAAGAAATTTGAGAGCA	1744	QY	2741	GGGCGGACTTCAGCTCCAGTGGCCACCCAGACCAAGATGAGAAAAACGAGAAACGATTAAT	2800	
DB	1750	AGGGGAACTTCAAGATATCAGATGTCGATTTGACCAACCAAGGCAAGAAATTTGAGAGCA	1809	DB	2830	GGGCGGACTTCAGCTCCAGTGGCCACCCAGACCAAGATGAGAAAAACGAGAAACGATTAAT	2889	
QY	1745	CAAAACAACTAGAGAGTTGAGAAATGCGCGGAATCACCCTCATCTACAGCACTAATTTAC	1800	QY	2801	GGGATGATGGGAGCCAGCCCT - TCTCTCACCTTCCAGTTCGCGGCGAGTTAAGGCGAG	2859	
DB	1810	CAAAACAACTAGAGAGTTGAGAAATGCGCGGAATCACCCTCATCTACAGCACTAATTTAC	1869	DB	2890	GGGATGATGGGAGCCAGCCCTTCTCTCACCGTTCCAGTTCGCGGCGAGTTAAGGCGAG	2949	
QY	1801	CAGGAATCTCAGCAAA - - -TGCTTGAAGACTTTATTCAGAAAAACAGATCTCAATGAC	1857	QY	2860	AGGTCGCTTTTACTCCAGCCAGGCGCACTGCTCTCCCTCTCTCTCTCTCTCTCTCTCTCT	2919	
DB	1870	ATGGAATCTCAGCAAAATGCTTTTGAAGAACTTTATTCAGAAAAACAGATCTCAATGAC	1929	DB	2950	AGGTCGCTTTTACTCCAGCCAGGCGCACTGCTCTCTCCCTCTCTCTCTCTCTCTCTCTCT	3009	

Qy	795	TAAATTACAAAAGGCACAGTCATTTGATGTGGCCAGTGTCCTCCACAGTGCAGAGTGGGC	854
Db	864	TAAATTACAAAAGGCACAGTCATTTGATGTGGCCAGTGTCCTCCACAGTGCAGAGTGGGC	923
Qy	855	TGTTCTCAGTCATCAAGATCGAAATACAGGCAATTAATCAATAGTCATGACAAAACATAT	914
Db	924	TGTTCTCTCAGTCATCAAGATCGAAATACAGGCAATTAATCAATAGTCATGACAAAACATAT	983
Qy	915	GAGTGGACACTTAAC-----AGGTCCCAAGCAAGAACTATTTCTTATGCAATC	962
Db	984	GAGTGGACACTTAACAGGTTCTGTTTAGGTCCCAAGCAAGAACTATTTCTTATGCAATC	1043
Qy	963	AAAGTTTACCACAGGCTCAGCTGGCTTCAATATGGAATCTTTTCTGACATTTGATCAAGATGG	1022
Db	1044	AAAGTTTACCACAGGCTCAGCTGGCTTCAATATGGAATCTTTTCTGACATTTGATCAAGATGG	1103
Qy	1023	AAAACCTTACAGCAGAGGAATTTATCTCGGCNATGCACTCATTTGATGTAGTACTATGCTGG	1082
Db	1104	AAAACCTTACAGCAGAGGAATTTATCTCGGCNATGCACTCATTTGATGTAGTACTATGCTGG	1163
Qy	1083	CCAACTACTGCACCTGTCCTGCCCTCCAGAAATACATTCACCTTCTTTTAGAAGAGTTTCG	1142
Db	1164	CCAACTACTGCCACTGTCCTGCCCTCCAGAAATACATTCACCTTCTTTTAGAAGAGTTTCG	1223
Qy	1143	ATCTGGCAGTGGTATATCTGTCTATPAGCTCAACATCTGTAGATCAGAGGCTACCAAGAGGA	1202
Db	1224	ATCTGGCAGTGGTATATCTGTCTATPAGCTCAACATCTGTAGATCAGAGGCTACCAAGAGGA	1283
Qy	1203	ACCAGTTTTAGAAAGATGAAACAAACAATTTAGAAAGAANAATTACCTCTACGTTTGNAGA	1262
Db	1284	ACCAGTTTTAGAAAGATGAAACAAACAATTTAGAAAGAANAATTACCTCTACGTTTGNAGA	1343
Qy	1263	TAAAGACGGGAGAACTTTGAACTGTGCAACTCTGGAACCTGGAGAAACGAAAGGCAAGCTCT	1322
Db	1344	TAGAAAGGGGAGAACTTTGAACTGTGCAACTCTGGAACCTGGAGAAACGAAAGGCAAGCTCT	1403
Qy	1323	CCTGGAAACAGCAGCGCAAGGAGCAGGAGCGCTGGCCACAGCTGGAGCGGGCGGAGCAAGGA	1382
Db	1404	CCTGGAAACAGCAGCGCAAGGAGCAGGAGCGCTGGCCACAGCTGGAGCGGGCGGAGCAAGGA	1463
Qy	1383	GAGGAAGGCGCTCAGCGCCAGGAGCAAGAGCGCAAAAGACAACCTGGAACCTGGAGGAGCA	1442
Db	1464	GAGGAAGGCGCTGAGCGCCAGGAGCAAGAGCGCAAAAGACAACCTGGAACCTGGAGGAGCA	1523
Qy	1443	ACTGGAAAAAGCAGCGGAGCTTAGAAACGGCAGAGAGAGGAGAGGAGGAGGAGGAGGAGGAGG	1502
Db	1524	ACTGGAAAAAGCAGCGGAGCTTAGAAACGGCAGAGAGAGGAGAGGAGGAGGAGGAGGAGGAGG	1583
Qy	1503	GAGCGGAGAGGCTGCAAAAACGGGAACTTTGAAAGGCAACGAACTTGAGTTGGGNAACGGAA	1562
Db	1584	GAGCGGAGAGGCTGCAAAAACGGGAACTTTGAAAGGCAACGAACTTGAGTTGGGNAACGGAA	1643
Qy	1563	TCGAAGGCAAGAACTACTTAAATCAAGAGAAACAAAGAGAAACAGAGGACATAGTTGTACTGAA	1622
Db	1644	TCGAAGGCAAGAACTACTTAAATCAAGAGAAACAAAGAGAAACAGAGGACATAGTTGTACTGAA	1703
Qy	1623	AGCAAGAAAAAGACTTTGGAAATTTGAAATTGAAAGCTCTTAAATGATAAAAAGCATCAACT	1682
Db	1704	AGCAAGAAAAAGACTTTGGAAATTTGAAATTGAAAGCTCTTAAATGATAAAAAGCATCAACT	1763
Qy	1683	AGAGGGAAACTTCAAGATATCAGATGTCGATTTGACCAACCCMAAGGCAAGAAATTGGAG	1742
Db	1764	AGAGGGAAACTTCAAGATATCAGATGTCGATTTGACCAACCCMAAGGCAAGAAATTGGAG	1823
Qy	1743	CACAAAACAAATCTAGAGAGTTTGAAATTTGCCAAATCACCCCATCTACAGCAACAAATTTACA	1802
Db	1824	CACAAAACAAATCTAGAGAGTTTGAAATTTGCCAAATCACCCCATCTACAGCAACAAATTTACA	1883
Qy	1803	GGAATCTCAGAAATGCTTGGAAGACTTATTTCCAGAAAAACAGATATCTCAATGACCAATT	1862
Db	1884	GGAATCTCAGAAATGCTTGGAAGACTTATTTCCAGAAAAACAGATATCTCAATGACCAATT	1943

QY	1863	AAAAAAGTTTCAGCAGAAACAAGTTTGGACAGAGATTCACCTTGTTTACACTTTAAAAGACGCTT	1922
DB	1944	AAAAAAGTTTCAGCAGAAACAAGTTTGGACAGAGATTCACCTTGTTTACACTTTAAAAGACGCTT	2003
QY	1923	AGAAAGCAAAAGAACTAGCTCGGCAGACACCTTACGAGACCAACTGCGATGAAAGTGGAGAAAGA	1982
DB	2004	AGAAAGCAAAAGAACTAGCTCGGCAGACACCTTACGAGACCAACTGCGATGAAAGTGGAGAAAGA	2063
QY	1983	AACCTAGATCAAAACTACAGGAGATTGATATTTTCAATAATACAGCTGAAAGAACTAAAGAGA	2042
DB	2064	AACCTAGATCAAAACTACAGGAGATTGATATTTTCAATAATACAGCTGAAAGAACTAAAGAGA	2123
QY	2043	AATACACAATAACGACAACCTCCAGAGCAAAAGTCCATGGAGGCTGAAACGACTGAAAAACA	2102
DB	2124	AATACACAATAACGACAACCTCCAGAGCAAAAGTCCATGGAGGCTGAAACGACTGAAAAACA	2183
QY	2103	GAAGAACAAGAAACGAAAGATCATAGAAATTTAGAAAAACAANAAGAGAAAGCCCAAGACG	2162
DB	2184	GAAGAACAAGAAACGAAAGATCATAGAAATTTAGAAAAACAANAAGAGAAAGCCCAAGACG	2243
QY	2163	AGCTCAGGAAAGGGAACAAGCAGTGGCTGGAGCATGTGCAGCAGGAGGACGAGCATCAGAG	2222
DB	2244	AGCTCAGGAAAGGGAACAAGCAGTGGCTGGAGCATGTGCAGCAGGAGGACGAGCATCAGAG	2303
QY	2223	ACCAAGAAAACTCCACAGCAGGAAAACTGAANAAGGAGGAGAGTGTCAANAAGAAAGGA	2282
DB	2304	ACCAAGAAAACTCCACAGCAGGAAAACTGAANAAGGAGGAGAGTGTCAANAAGAAAGGA	2363
QY	2283	TGGCGAGGAAAAAGGCAACACAGAAAGCAACAACAAGCTGGGTGGCTTTTCATCAACA	2342
DB	2364	TGGCGAGGAAAAAGGCAACACAGAAAGCAACAACAAGCTGGGTGGCTTTTCATCAACA	2423
QY	2343	CCAGAACACAGCTAAGCCAGCTCTCCAGGACACCTCGTCCACTGCAGAAAAAGGTCCACT	2402
DB	2424	CCAGAACACAGCTAAGCCAGCTCTCCAGGACACCTCGTCCACTGCAGAAAAAGGTCCACT	2483
QY	2403	TACCATTCTCTGCACAGGAAAAATGAAAGTGGTGTATTATCCCGGGCACTGTACCCCTTTGA	2462
DB	2484	TACCATTCTCTGCACAGGAAAAATGAAAGTGGTGTATTATCCCGGGCACTGTACCCCTTTGA	2543
QY	2463	ATCCAGAAAGCCATGATGAATATCATATCCAGCCAGGAGACATAGTCATGGTGGATCAAG	2522
DB	2544	ATCCAGAAAGCCATGATGAATATCATATCCAGCCAGGAGACATAGTCATGGTGGATCAAG	2603
QY	2523	CCAAACTGGAGAACCCCGCTGGCTTGGAGAGAAATTAANAAGGAAAGACAGGGTGGTTCCC	2582
DB	2604	CCAAACTGGAGAACCCCGCTGGCTTGGAGAGAAATTAANAAGGAAAGACAGGGTGGTTCCC	2663
QY	2583	TGCAAACTATTCAGAGAAAAATCCACAGAAAAATGAGGTTCCCGCTCCAGTGAAACCAAGTGAC	2642
DB	2664	TGCAAACTATTCAGAGAAAAATCCACAGAAAAATGAGGTTCCCGCTCCAGTGAAACCAAGTGAC	2723
QY	2643	TGATTTCAACATCTGCCCTCCCGCTCCCGCTTGGCTGGAGACCCCGCCCTTTGGC	2702
DB	2724	TGATTTCAACATCTGCCCTCCCGCTCCCGCTTGGCTGGAGACCCCGCCCTTTGGC	2783
QY	2703	AGTAACCTCTTCAGAGCCCTCCACGACCCCTAATAACTGGGCGCATTTACGTCTCCACGTG	2762
DB	2784	AGTAACCTCTTCAGAGCCCTCCACGACCCCTAATAACTGGGCGCATTTACGTCTCCACGTG	2843
QY	2763	GCCACACGACACGATGAGAAACAGNAACGGATACTGGGATGCTATGGGACGCCAGGCC	2822
DB	2844	GCCACACGACACGATGAGAAACAGNAACGGATACTGGGATGCTATGGGACGCCAGGCC	2903
QY	2823	CTCTCTCAACCGTTCCAAAGTCGCGGCCAGTTTAAAGGACAGAGTCCGCCCTTTACTCCAGCCAC	2882
DB	2904	CTCTCTCAACCGTTCCAAAGTCGCGGCCAGTTTAAAGGACAGAGTCCGCCCTTTACTCCAGCCAC	2963
QY	2883	GGCCACTGGCTCTCTCCCGCTCTCTCTGTGTAGGCCAGGGTGA	2944
DB	2964	GGCCACTGGCTCTCTCCCGCTCTCTCTGTGTAGGCCAGGGTGA	3005

RESULT 6			US-09-764-875-88		
			; Sequence 88, Application US/09764875		
			; Publication No. US20040018969A1		
			; GENERAL INFORMATION:		
			; APPLICANT: Rosen et al.		
			; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies		
			; FILE REFERENCE: PJ202		
			; CURRENT APPLICATION NUMBER: US/09/764,875		
			; CURRENT FILING DATE: 2001-01-17		
			; Prior application data removed - consult PALM or file wrapper		
			; NUMBER OF SEQ ID NOS: 1249		
			; SOFTWARE: PatentIn Ver. 2.0		
			; SEQ ID NO 88		
			; LENGTH: 3319		
			; TYPE: DNA		
			; ORGANISM: Homo sapiens		
			US-09-764-875-88		
			Query Match 54.7%; Score 2843.2; DB 3; Length 3319;		
			Best Local Similarity 99.5%; Pred. No. 0;		
			Matches 2867; Conservative 0; Mismatches 3; Indels 12; Gaps 1;		
QY	55	GACGGACAGAGGCGGGGGGATGGTGGCGGGCTGGCGGCTCTGGGCTCCCTGCCAG	114		
DB	6	GAGGGACAGAGGCGGGGGGATGGTGGCGGGCTGGCGGCTCTGGGCTCCCTGCCAG	65		
QY	115	CGGCGGCTGAGCGGCACTGATTTGTCCCTGGGCGGCGAGCGCGACCCCGCGGAGTGA	174		
DB	66	CGGCGGCTGAGCGGCACTGATTTGTCCCTGGGCGGCGAGCGCGACCCCGCGGAGTGA	125		
QY	175	GGCGTGCATTTAGCAAGGTAAGTAACAGAACCATGGCTCAGTTTCCACACCTTTTGGT	234		
DB	126	GGCGTGCATTTAGCAAGGTAAGTAACAGAACCATGGCTCAGTTTCCACACCTTTTGGT	185		
QY	235	GGCAGCTGGATATCTGGGCGTAATCTGTAGAGAAAGAGCGAGCATGATCAGCAGTTC	294		
DB	186	GGCAGCTGGATATCTGGGCGTAATCTGTAGAGAAAGAGCGAGCATGATCAGCAGTTC	245		
QY	295	CATAGTTTAAAGCAATATCTGGATTCATTTACTGGTGATCAAGCTAGAACTTTTTTTT	354		
DB	246	CATAGTTTAAAGCAATATCTGGATTCATTTACTGGTGATCAAGCTAGAACTTTTTTTT	305		
QY	355	CAATCTGGTTACTCAACTGTTTGTAGCAAGATATGGCACTAGCTGATGAATAAT	414		
DB	306	CAATCTGGTTACTCAACTGTTTGTAGCAAGATATGGCACTAGCTGATGAATAAT	365		
QY	415	GATGGAGATGATCAAGTGGAGTTTCCATAGCTATGAACTTATCAAACTGAAGCTA	474		
DB	366	GATGGAGATGATCAAGTGGAGTTTCCATAGCTATGAACTTATCAAACTGAAGCTA	425		
QY	475	CAAGGATATCAGCTACCCCTCTGCACCTTCCCTCTGTCTATGAACAGCAACAGTTGCTATT	534		
DB	426	CAAGGATATCAGCTACCCCTCTGCACCTTCCCTCTGTCTATGAACAGCAACAGTTGCTATT	485		
QY	535	TCTAGCCACAGCATTTGGTATGGGAGGTATGCCAGCATGCCACCGCTTACAGCTGTT	594		
DB	486	TCTAGCCACAGCATTTGGTATGGGAGGTATGCCAGCATGCCACCGCTTACAGCTGTT	545		
QY	595	GTCTCCAGTCCAAATGGGATCCATTCAGTTGTTGGAGTTGTTCCAACTAGTATCTTCT	654		
DB	546	GTCTCCAGTCCAAATGGGATCCATTCAGTTGTTGGAGTTGTTCCAACTAGTATCTTCT	605		
QY	655	GTTCACACAGCAGCTGTGCCCTCTGGCTAACCGGGCTCCCTGTTATACAACTCTG	714		
DB	606	GTTCACACAGCAGCTGTGCCCTCTGGCTAACCGGGCTCCCTGTTATACAACTCTG	665		
QY	715	CTGCAATTTGCTATCTCTGAGCCACAATTTGCAAGAGTTCTTCTTTAGTAGTCTGGT	774		
DB	666	CTGCAATTTGCTATCTCTGAGCCACAATTTGCAAGAGTTCTTCTTTAGTAGTCTGGT	725		
QY	775	CCAGGCTCAACACTAAACCTAAATTTACAAAGGCACAGTCATTTGATGTGGCCAGTGC	834		

DB	726	CCAGGGTCACAACTAAACACTAAATTTACAAAAGGCACAGTCATTTGATGGCCAGTGC	785
QY	835	CCACCAGTGGCAGAGTGGGCTGTTCTCAGTCATCAAGACTGAAATACAGGCAATTTATTC	894
DB	786	CCACCAGTGGCAGAGTGGGCTGTTCTCAGTCATCAAGACTGAAATACAGGCAATTTATTC	845
QY	895	AATAGTCATGACAAAACTATGATGGACACTTTAAC-----AGTCTCCCAAGCA	942
DB	846	AATAGTCATGACAAAACTATGATGGACACTTTAACAGAGTTCTCTGTTTAGTCTCCCAAGCA	905
QY	943	AGAACTATTTCTTATGAGCTCAAGTTTACACAGGCTCAGCTGGCTTCAATATGGAATCTT	1002
DB	906	AGAACTATTTCTTATGAGCTCAAGTTTACACAGGCTCAGCTGGCTTCAATATGGAATCTT	965
QY	1003	TCTGACATTTGATCAAGTGGAAAACTTACAGCAGAGGAAATTTATCTGGCAATGCACTTC	1062
DB	966	TCTGACATTTGATCAAGTGGAAAACTTACAGCAGAGGAAATTTATCTGGCAATGCACTTC	1025
QY	1063	ATTGATGTAGCTATGTCTGGCCAAACCACTGCCACCTGTCTGCTCCAGAAATCATTTCCA	1122
DB	1026	ATTGATGTAGCTATGTCTGGCCAAACCACTGCCACCTGTCTGCTCCAGAAATCATTTCCA	1085
QY	1123	CTTCTCTTTTAGAAGAGTTTCGATCTGCGAGTGGTATATCTGTCTATAGCTCAACATCTGA	1182
DB	1086	CTTCTCTTTTAGAAGAGTTTCGATCTGCGAGTGGTATATCTGTCTATAGCTCAACATCTGA	1145
QY	1183	GATCAGAGGCTACACAGAGAACCCAGTTTGTAGAAAGTGAACAACAACAAATTTAGAAAAAGAA	1242
DB	1146	GATCAGAGGCTACACAGAGAACCCAGTTTGTAGAAAGTGAACAACAACAAATTTAGAAAAAGAA	1205
QY	1243	TTACTCTGTAACTGTTTGAAGATAAGAACCGGGAGAACTTTGAACTGGCAACCTGGAACTG	1302
DB	1206	TTACTCTGTAACTGTTTGAAGATAAGAACCGGGAGAACTTTGAACTGGCAACCTGGAACTG	1265
QY	1303	GAGAAAACGAAAGCAAGCTCTCTGGAACAGCAGCAGCAAGAGCAGGAGCCCTGGCCAG	1362
DB	1266	GAGAAAACGAAAGCAAGCTCTCTGGAACAGCAGCAGCAAGAGCAGGAGCCCTGGCCAG	1325
QY	1363	CTGAGCGGGCGGAGCAGGAGAGGAGGAGCGTGGCGCCAGGAGCAAGAGCGCAAAAGA	1422
DB	1326	CTGAGCGGGCGGAGCAGGAGAGGAGGAGCGTGGCGCCAGGAGCAAGAGCGCAAAAGA	1385
QY	1423	CAACTGGAACCTGGAGAGCAACTCTGGAAGACAGCGGAGCTAGAAACGCGAGAGAGAG	1482
DB	1386	CAACTGGAACCTGGAGAGCAACTCTGGAAGACAGCGGAGCTAGAAACGCGAGAGAGAG	1445
QY	1483	GAGAGGAGGAAAGAAATTTGAGAGCGGAGAGGCTGCAAAAACGGGAACTTTGAAAGCAACGA	1542
DB	1446	GAGAGGAGGAAAGAAATTTGAGAGCGGAGAGGCTGCAAAAACGGGAACTTTGAAAGCAACGA	1505
QY	1543	CAACTTGGTGGGAAACGGAATCGAAGCAAGAACTACTTAATCAAGAAACAAAGAAACAA	1602
DB	1506	CAACTTGGTGGGAAACGGAATCGAAGCAAGAACTACTTAATCAAGAAACAAAGAAACAA	1565
QY	1603	GAGGACATAGTTGTTACTGAAAGCAAAAGAAAGACTTTTGAATTTGAATAGAGCTCTA	1662
DB	1566	GAGGACATAGTTGTTACTGAAAGCAAAAGAAAGACTTTTGAATTTGAATAGAGCTCTA	1625
QY	1663	AATGATAAAAAGCATCAACTAGAGGAAACTTCAAGATATCAGATCTCGATTGACCAACC	1722
DB	1626	AATGATAAAAAGCATCAACTAGAGGAAACTTCAAGATATCAGATCTCGATTGACCAACC	1685
QY	1723	CAAGGCAAGAAATTTGAGAGCACAACAAATCTAGAGAGTTGAGAAATTCGCGAAATCAAC	1782
DB	1686	CAAGGCAAGAAATTTGAGAGCACAACAAATCTAGAGAGTTGAGAAATTCGCGAAATCAAC	1745
QY	1783	CATCTACAGCAACAAATTTACAGGAACTCTCAGCAATGCTTGGAGAGCTTTATTTCCAGAAAA	1842
DB	1746	CATCTACAGCAACAAATTTACAGGAACTCTCAGCAATGCTTGGAGAGCTTTATTTCCAGAAAA	1805
QY	1843	CAGATCTCAATGACCAATTTAAACAAAGTTTCAGAGAACAGTTTGGCAGAGATTTCACTT	1902
DB	1806	CAGATCTCAATGACCAATTTAAACAAAGTTTCAGAGAACAGTTTGGCAGAGATTTCACTT	1865

533 TCCTCAAGAGAGCCACTATCCCATATCACTGCCAGCA-GGATGATGGAGATGCAGCCT 591
3739 TGATCATGTGACTTCCAGCATGATCACTATCGCTTCTGAGTAGAAGAACTCACTGGAG 3798
592 TGATCATGTGACTTCCAGCATGATCACTATCGCTTCTGAGTAGAAGAACTCACTGGAG 651
3799 AGAGTTTACCTCATTTTACCTTAGTGTGATGATGCGAATGTTGAGTTTATTAATTGC 3858
652 AGCAKTTTACCTCATTTTACCTTAGTGTGATGATGCGAATGTTGAGTTTATTAATTGC 711
3859 AGAGTAGGAGCAAAATTTACAAACACACAGAGGTAGTGGTCCCTTTTGTGGCTTTCCT 3918
712 AGAGTAGGAGCAAAATTTACAAACACACAGAGGTAGTGGTCCCTTTTGTGGCTTTCCT 771
3919 AGTTACTCAAAATGACTTCCCCACCTTTGACAGGTGCTTCAATAGTGTATAAATTA 3978
772 AGTTACTCAAAATGACTTCCCCACCTTTGACAGGTGCTTCAATAGTGTATAAATTA 831
3979 TTTTAAATATATATTTTACCTTTTAAATAAATAAATAAATAAATAAATAAATAAATAA 4038
832 TTTTAAATATATATTTTACCTTTTAAATAAATAAATAAATAAATAAATAAATAAATAA 891
4039 TTTGGTTTGCAGAGAGCCACTATCAAGGAATGCTGCATGCTGCTATTAAATAATTTGTT 4098
892 TTTGGTTTGCAGAGAGCCACTATCAAGGAATGCTGCATGCTGCTATTAAATAATTTGTT 951
4099 CAAATGCTCATAAATCTGAGACTTGTATGATTTTTTCAATTTGTCAGTGTACCAACTA 4158
952 CAAATGCTCATAAATCTGAGACTTGTATGATTTTTTCAATTTGTCAGTGTACCAACTA 1011
4159 AATGCTGCAAGTTGGGGCTTTTCCCTTTACATAGAGTGCAGAGGATTCAGTATCT 4218
1012 AATG-TGCAGTTGGGGCTTTTCCCTTTACATAGAGTGCAGAGGATTCAGTATCT 1070
4219 CTGTTTAAAGAGTATAGATGAGCCCAATTAAGCGAAGTGTGCTTGTGTTTG 4278
1071 CTGTTTAAAGAGTATAGATGAGCCCAATTAAGCGAAGTGTGCTTGTGTTTG 1130
4279 TGTATCAGCTGTACCTTGTGAGCATGTAATACATCTGTACATAAGAAATTTAGTCTTT 4338
1131 TGTATCAGCTGTACCTTGTGAGCATGTAATACATCTGTACATAAGAAATTTAGTCTTT 1190
4339 CCATGGCAAGCTATTAACCTTGTACGATGCTTAATCAATATGCAATTAATTTATTTG 4398
1191 CCATGGCAAGCTATTAACCTTGTACGATGCTTAATCAATATGCAATTAATTTATTTG 1250
4399 CAACAGTGACCTTGTAGCCATGAGAAAGCACTCTGTTTGTGTTGCTGCTCAGATTT 4458
1251 C-ACAGTGACCTTGTAGCCATGAGAAAGCACTCTGTTTGTGTTGCTGCTCAGATTT 1309
4459 ATCTGGTTGAGTTGGTGTGTTTGGGGTTTTTAAATTTTGGGTTTGCATAGCATPAA 4518
1310 ATCTGGTTGAGTTGGTGTGTTTGGGGTTTTTAAATTTTGGGTTTGCATAGCATPAA 1369
4519 ATCAGTAGACAACCACTAGGTCGTTACGATCAAGATATCAGAGTCTCTTTTGTAGT 4578
1370 ATCAGTAGACAACCACTAGGTCGTTACGATCAAGATATCAGAGTCTCTTTTGTAGT 1429
4579 CTCGTTTACATGAATTTTATTCAGTTTACTTTTTCATGGAATGACCTTTTGAACAGT 4638
1430 CTCGTTTACATGAATTTTATTCAGTTTACTTTTTCATGGAATGACCTTTTGAACAGT 1489
4639 AATTTTCTTGAAGAAAGATGTATAGAGTCTCCCTGCAATTAATTTTCAATGTTTAC 4698
1490 AATTTTCTTGAAGAAAGATGTATAGAGTCTCCCTGCAATTAATTTTCAATGTTTAC 1549
4699 ATTTTAACTPAGGACTGTGAAATTTCTACAGATTAATGAAATGAGGCTCATGGTCCG 4758
1550 ATTTTAACTA-GACTGTGAAATTTCTACAGATTAATGAAATGAGGCTCATGGTCCG 1608
4759 TTTGTGTTAGATATGCTGAGGCTGTTGTTGCTTTTAAACACTAGTTGGAG 4818
1609 TTTGTGTTAGATATGCTGAGGCTGTTGTTGCTTTTAAACACTAGTTGGAG 1668

4819 CTCTCAATAAAAAATGCCTGCTCTCACAGCACAGAAAAATGGGGGAGGAGCCTCAAGC 4878
1669 CTCTCAATAAAAAATGCCTGCTCTCACAGCACAGAAAAATGGGGGAGGAGCCTCAAGC 1728
4879 ACAATCTAGCTGTCTCTCTAAAGACTCTGTAATGCTCAATCCCTTTCCTCCGGCG 4938
1729 ACAATCTAGCTGTCTCTCTAAAGACTCTGTAATGCTCAATCCCTTTCCTCCGGCG 1788
4939 CTGTGGGAGGCTGTGCTGCTGTAGAGTCCCTTTTCTTTTCAAAATGCTGAGAG 4998
1789 CTGTGGGAGGCTGTGCTGCTGTAGAGTCCCTTTTCTTTTCAAAATGCTGAGAG 1848
4999 AGAGAGGAGCTTTCTCTCTTGTTCAGTTTGCATTTTCAAGTATTTTACGGATATGAATGTA 5058
1849 AGAGAGGAGCTTTCTCTCTTGTTCAGTTTGCATTTTCAAGTATTTTACGGATATGAATGTA 1908
5059 AATATATAAATAATAAATCAACCTGAGGATTTAAACAATGTAAACAACCTTTTGAATTAGTT 5118
1909 AATATATAAATAATAAATCAACCTGAGGATTTAAACAATGTAAACAACCTTTTGAATTAGTT 1967
5119 CCGAGTATAGATAAATAAATTTTAAACAAGTAAAAAAGTAAAAAAGTAAAAAAGTAAAA 5178
1968 CCGTGTATAGATAAATAAATTTTAAACAAGTAAAAAAGTAAAAAAGTAAAAAAGTAAAA 2027
5179 AAAAAA 5184
2028 AAAAAA 2033

RESULT 8
US-10-450-763-20566
; Sequence 20566, Application US/10450763
; Publication No. US20050196754A1
; GENERAL INFORMATION:
; APPLICANT: Hyseq, Inc
; TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES
; FILE REFERENCE: 790CIP3/US
; CURRENT APPLICATION NUMBER: US/10/450,763
; CURRENT FILING DATE: 2003-06-11
; PRIOR APPLICATION NUMBER: PCT/US01/08631
; PRIOR FILING DATE: 2001-03-30
; PRIOR APPLICATION NUMBER: 09/540,217
; PRIOR FILING DATE: 2000-03-31
; PRIOR APPLICATION NUMBER: 09/649,167
; PRIOR FILING DATE: 2000-08-23
; NUMBER OF SEQ ID NOS: 60736
; SOFTWARE: Custom
; SEQ ID NO 20566
; LENGTH: 2874
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SIMILAR
; LOCATION: (1)..(1545)
; OTHER INFORMATION: 100% homologous to Homo sapiens Human SH3D1A
; OTHER INFORMATION: protein, accession number Y32156, Smith-Waterman Score=2669.
US-10-450-763-20566

Query Match 31.4%; Score 1630; DB 9; Length 2874;
Best Local Similarity 87.5%; Pred. No. 0;
Matches 1862; Conservative 0; Mismatches 185; Indels 81; Gaps 4;
Qy 1513 GCTGCAAAACGGGAACTTGAAGCAACGACCACTTCAAGTGGGAACCGAATCGAAGGCAA 1572
Db 1 GCTGCAAAACGGGAACTTGAAGCAACGACCACTTCAAGTGGGAACCGAATCGAAGGCAA 60
Qy 1573 GAACCTACTTAATCAAGAAACAAAGAGACACATAGTTGTACTGAAACGAAGAAA 1632
Db 61 GAACCTACTTAATCAAGAAACAAAGAGACACATAGTTGTACTGAAACGAAGAAA 120
Qy 1633 AAGACTTTGGAAATTTGAATTAAGAGCTCTAAATGATAAAAAGCATCAACTAGAGGAAA 1692


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; APPLICANT: Takahasi, Biki
; TITLE OF INVENTION: Method of Testing For Allergic Diseases
; FILE REFERENCE: SHIMIZU-07907
; CURRENT APPLICATION NUMBER: US/10/398,885A
; CURRENT FILING DATE: 2003-08-11
; PRIOR APPLICATION NUMBER: PCT/Jp01/08937
; PRIOR FILING DATE: 2001-10-11
; PRIOR APPLICATION NUMBER: JP 2000-314093
; PRIOR FILING DATE: 2000-10-13
; NUMBER OF SEQ ID NOS: 16
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 15
; TYPE: DNA
; LENGTH: 5828
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Synthetic
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (7)..(5052)
; OTHER INFORMATION:
; US-10-398-885A-15

Query Match          10.6%; Score 550; DB 7; Length 5828;
Best Local Similarity 50.7%; Pred. No. 1.4e-116;
Matches 1811; Conservative 0; Mismatches 1595; Indels 164; Gaps 14;

QY      184  TACCAAGGTAAAGTAACAGAACCCNTGGCTCAGTTTTCCACACACTTTTGGTGCGACGCTG 243
Db      184  TATCAAGGAAAACTCAGGACCATGATGGCTCAGTTTTCCACAGCTATGAATGGAGGGCCA 78

QY      244  GATATCTGGGCCATAACTGTAGAGGAAGAGCGAAGCATGATCAGCAGTTCCCATAGTTTA 303
Db      244  AACATGTGGGCTATTACCTCTGAAGACGTAAGCATGACGCGAGTTTGATAACCTC 138

QY      304  AAGCCAAATCTCGGATTCATTACTGGTGATCAAGCTAGAACTTTTTTTTTCAATCTGGG 363
Db      304  AAACCTTCAGGAGGTTACATAACAGGTGATCAAGCAGGTAAATTTTCTCAATCAAGT 198

QY      364  TTACCTCAACTGTGTTTAGCACAGATATGGSCATCTAGCTGACATGAATAATGATGGAAGA 423
Db      364  CTGCGGCCCCCTGTTTTAGCTGAATATGGCTTTATCAGACCTTAACNAGATGGGAG 258

QY      424  ATGGATCAAGTGGAGTTTTCATAGCTATGAAACTTATCAAACTGAAGCTACAGGATAT 483
Db      424  ATGGATCAGCAAGAGTTTCTCCATAGCTATGAAACTCATCAAACTGAAGCTTCAAGGCCAA 318

QY      484  CAGTACCTCTGCACACTTCCCTCTCATGAAACAGCAACC---AGTTGCTATTTCCTAGC 540
Db      484  CAGTTGCTGTGGTTCCTCCCTCTTATATGAAGCAACCCCTATGTTTTCTCCATTAATT 378

QY      541  GCACCAAGATTTGGTATGGGAGGTATGCCACAGCATGCCACGCTTACAGCTGTGTCTCCA 600
Db      541  TCTGCTCGTTTTGGAATGGGAAGCATGCCCAATCTGTCCATTCTCAGCCATTGCTCCA 438

QY      601  GTGCCAATGGGATC-----CAITTCAGTTGTTGGA 630
Db      601  GCTGCACCTATAACATCATTTGTTCTCTCGCATTCAGGGACCAACCTTCTCCCTTAAATG 498

QY      631  ATGTCCTCAACCCTAGTATCTTCGTGCCACAGCAGCTGTGCCCCCTGGCTAACCGGG 690
Db      631  ATGCCCCATCCCTCTAGTGCCCTTCGTGTAGCAATCATCATTTACCAAAATGGAAACCCAGT 558

QY      691  GCTCCCCCTGTATACAACTCTGCCTGCATTTGCTCATCTCGACGCCACATGTCACAAG 750
Db      691  CTCATTGAGCCCTTTACCCATTCTTATTTCTTCAACATTTGCCCTCATGGGTCATCTTAT 618

QY      751  AGTTCTTCTTTAGTAGATCTGGTCCAGGGTCACAACTAAACACTAAATACAAAGGCA 810
Db      751  AGTCTGATGATGGGAGGATTTGGAGGTCTAGTATACAGAAGCCGAGTCTCTGATTGAT 678

QY      811  CAGTCATTTGATGTGGCCAGTGTGCCACGAG----- 841

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QY 1918 GCTTTAGAACAAAAGAACTAGCTCGGACGACCTACGAGACCAACTGATGAAGTGGAG 1977
Db 1789 AAATCATTAAGAAAAGAAATTAATGCCAAAGACTTAAAGAAACAGTTAGATCTCTTGA 1848
QY 1978 AAAGAACTAGATCAAAACTACAGGAGATTGATATTTTCAATAATCAGCTGAAGGACTA 2037
Db 1849 AAAGAAACTGCACTTAAGCTGTAGAAATGGATTCCTTTAAACAATCAACTAAAGGAACTG 1908
QY 2038 AGAGAAATACACAATAAGCAACAACCTCCAGAAAGCAAAAGTCCATGGAGGCTGAAACGACTG 2097
Db 1909 AGAGAAACCTTACAACACAGCAGTGTAGCCCTTGAACACTTTATAGATCAAACTGTAC 1968
QY 2098 AAACAGAAAGACAAAGAACGAAAGATCATAGAAAT-----AGAAAAACAAAAGAA 2148
Db 1969 AAGTTGAAGGAAATTCGAAAGGAAAGATTTAGAACTAATGCAGAAAAAGAAACTTAGAAT 2028
QY 2149 GAAGCCAAAGCAGCTCAGGAAAGGACAGCAGTGCCTGGAGCATGTGCAGCAGGAG 2208
Db 2029 GAGGCTGCAAGGAAAGCAAGCAAGGAAAGAAAGAAACTTATGAAAGAAAAATCTTAGAAAG 2088
QY 2209 GACGACATCAGAGACCAAGAAAACTCCACGAAGAGGAAAAACTGAAAAAGGGAGAGCT 2268
Db 2089 GAGGAGAGAAAAACAAAGCGACTCCAGGAGAAAAACACAGAAAAAATTCAGAA 2148
QY 2269 GTCAAAAGAAAGATGGCGAGGAAAAAGGAAACAGGAAG-CACAAGACAAGCTGGGTG 2327
Db 2149 GAGGAACGGAAGCTGAGGAGAAACAACGTAAGGATAAGGATACTTTTGAAGCTGAGGAG 2208
QY 2328 GCTTTTCCATCAACACCAAGACCAGCTAAGCCAGCTGCCAGGACCCCTGCTCCACTGC 2387
Db 2209 AAAAAACGTGAGACAGCTAGTGTGTTTGGTGAATATATAGAGCATATATACCCCTTTGA 2268
QY 2388 AGAAAAAGGTCCTACCTTACCATTCTGCACAGGAAAAATGTAAAGTGGTGTATTACCGGC 2447
Db 2269 AGGACCATGATGAGATGAGTTTAAATCTGGAGATATTAATTCAGGTTCATGAAAAACC 2328
QY 2448 ACTGATCCCTTTGAATCCAGAAAGCCATGATGAATCATATCCAGCCAGGAGACATAGT 2507
Db 2329 GTAGGAGAACCTGGTTGGCTTTATGTTAGTATTTTCAAGGAAATTTTGGCTGGTTTCATGC 2388
QY 2508 CATGGTGGATG-----AAGCCAACTGGAGAACCCGGCTGGCTTGGAGGAA 2556
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QY 2557 TTAAGAGGAAAGACAGGGTGGTTCCCTGCAAACT---ATGCAGAGAAAAATCCAGAAAAAT 2613
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QY 2614 GAGGTTCCCGCTCCAGTGAACCAAGTGACTGATTCAAACATCTGCCCTGCCCCCAAACTG 2673
Db 2509 AATCAACCAAGCATCAGTCACTGATTAATCAAAATGTATCTTTTCAAAACCTAACTGTAAT 2568
QY 2674 GCCTTCGTTGAGACCCCGCCCTTTGGCAGTAACTCTTCAGAGCCCTCCAGACCCCT 2733
Db 2569 ACATCATGCGAGAAAAAATCAGCCTTCACTCGAACTGTGTCCCTGGATCTGTATCACCT 2628
QY 2734 AATAACTGG-----GCCGACTTCAGCTCCACGTCGCGCCACACGACGAAATGAGAAA 2784
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QY 2785 CCAGAAACCGATAACTGGGATGATGGGAGCCAGCCCTCTCTCAACGTTCCAAAGTGGC 2844
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QY 2845 GGCCGTTAAGGCAGAGGTCGCGCTTTACTCCAGCCACGSCCACTGGCTCCCTCCCGTCT 2904
Db 2749 CAAGAAAAATTTGGTGTGGGAGGTGCATGGAGGAAGAGGATGGTTTCCCAAAATCTTAT 2808
QY 2905 CCTGTGCTAGGCGAGGGTGAAAAAGGTGGAGGGGCTACAAGCTCAAGCCCTTATCTCTGG 2964
Db 2809 GTCAAGATCATTTCTCGGAGTGAAGTAAAAACGGGAGAACCCAGAAAGCTTTGTATGACGT 2868
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QY 2965 AGAGCCAAAAAGACCAACCACTTAAATTTTAAACAAAATATGTCTATCACCGTCTCGGAA 3024
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QY 3025 CAGCAAGACATGTGTGGTGTGGAGAGCTTCAAGGTCTAGAAGGGTGGTTCCCCAAGTCT 3084
Db 2929 C-CATATTCAGTGTGGAACTCGGAGATTTCACITTTACAGAGGTGAAGAAATATTGGT 2987
QY 3085 TACGTGAAACTCATTTTCAGGGCCCATAGGAAGTCTACAAGCATGGATTCTGGTCTTCA 3144
Db 2988 GACCCAGAAAAGATGAGAGTGTGGACAGGAAGTATTGGAGATAGAGTGGAAATTTTTCC 3047
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Db 3048 ATCAAACTATGTCAAAACCAAAAGGATCAAGAGAGTTTGGGAGTGTAGCAAGCTCTGGAGC 3107
QY 3197 TTTCCGGAGAA-----GAAATTTGCCAGGTATTGCTCTATACACCGCCACCGGCCCGA 3251
Db 3108 ATCAATTAATAAAACCTGAGATTGCTCAGGTAACTTCAGCATATGTTGCTTCTGGTCTGA 3167
QY 3252 GCAGTCACTCTCGCCCTGTGTCAGTCAATTTTGTATCCGAAAAAGAACCCAGGTGGATG 3311
Db 3168 ACAACTTAGCCTTGCACCAGGACAGTTAATATTAATTTCTAAAGAAAAATACAAGTGGGTG 3227
QY 3312 GTGGGAAGGAGCTGCAAGCACGCTGGGAAAAAGCCAGATAGGCTGTTCCCGAGTAA 3371
Db 3228 GTGGCAAGGAGAGTTACAGGCCAGGAAAAAGGCAAGAAAGGATGTTTCTCTGCCAG 3287
QY 3372 TTATGTAAGCTTCTTAAGCCCTGGGACGAGCAAAATCACTCCAAACAGAGCCACCTAAGTC 3431
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QY 3432 AACACATAGCGGAGTGTGCCAGGTGATTTGGGATGTACGACTACACCGGCGAGATGA 3491
Db 3327 ACCTGCCCTTCATCTCTGTATGTGTCAGGTGATGCTATGTATGATGATGAGCAAAATGA 3386
QY 3492 CGATCAGCTGGCCTTCAACAAGGGCCAGATCATCAACGTCCTCAACAAGGAGGACCTGA 3551
Db 3387 AGATGAGCTCAGTTTCTCAAGGGCAACTCATTAATGTTATGAACAAGATGATCTCTGA 3446
QY 3552 CTGTTGGAAGAGAGTCAATGGACAAGTGGGCTCTTCCCATCCAATTTATGTGAAGCT 3611
Db 3447 TTGTTGGCAGGAGAGATCAACGGGTGACTGCTCTTCTTCTTCAAACTAGCTTAAGAT 3506
QY 3612 GACCAAGACATGGACCCCAAGCCAGCAATG 3641
Db 3507 GACGACAGACTCAGATCCAAGTCAACAGTG 3536
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RESULT 10

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US-09-884-441-72
; Sequence 72, Application US/09884441
; Patent No. US20020119158A1
; GENERAL INFORMATION:
; APPLICANT: Algate, Paul A.
; APPLICANT: Carcer, Darick
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
; FILE REFERENCE: 210121.462C7
; CURRENT APPLICATION NUMBER: US/09/884,441
; CURRENT FILING DATE: 2001-06-18
; NUMBER OF SEQ ID NOS: 489
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 72
; LENGTH: 2017
; TYPE: DNA
; ORGANISM: Homo sapien
US-09-884-441-72
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Query Match 9.8%; Score 507.8; DB 3; Length 2017;
Best Local Similarity 55.8%; Pred. No. 4.9e-107;
Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5;

QY	175	GGCGTCGATTAGCAAGGTAAGTAACAGAACCACTGGCTCAGTTTCCACACCTTTGGT	234
Db	9	GGCTGAGAGCTGCAGAAAGAGTCAGGATCATGATGGCTCAGTTTCCACAGGATGAAT	68
QY	235	GGCAGCGCTGGATATCTGGGCCATAACTGTAGAGGAAGAGCGAAGCATGATCAGCAGTTTC	294
Db	69	GGAGGCCAATATGTGGGCTATTACATCTGAAGACGCTACTAAGCATGATAAAGCTTT	128
QY	295	CATAGTTTAAAGCCAAATATCTGGATTCAATCTGTGGTGATCAAGCTAGAAACTTTTTTTT	354
Db	129	GATAACCTCAAACTTTCAGGAGGTTACATAACAGGTGATCAAGCCGCTACTTTTTTCCTA	188
QY	355	CAATCTGGGTTACCTCAACTGTTTTAGCACAGATATGGCACTAGCTGACATGAATAAT	414
Db	189	CAGTCAGGCTGCGGCCCGGTTTTAGCTGAATAATGGCCCTTATCAGATCTGAACAAG	248
QY	415	GATGGAAGATGATCAAGTGGAGTTTTCATAGCTATGAAACTTATCAAACTGAAGCTA	474
Db	249	GATGGGAAGATGGACCAAGAGTTCTCTATAGCTATGAACTCATCAAGTTAAAGTTG	308
QY	475	CAAGGATATCAGTACCTCTGCACTTCCCTCTGTCTATGAAACAGCAACG---AGTTGCT	531
Db	309	CAGGGCCAAAGCTGCTGTAGTCTCTCCCTCTATCATGAAACAGCCCTATGTTCTCT	368
QY	532	ATTTCTAGCCACCAAGATTTGGTATGGAGGTATCGCCAGCATGCCACCGCTTACAGCT	591
Db	369	CCACTAATCTCTGCTCGTTTTGGGATGGGAAGCATGCCCAATCTGTCCATTCAAGCCA	428
QY	592	GTTCCTCCAGTGCCAATGGGATC-----CAAT	618
Db	429	TTGCCCTCCAGTTGCCTATAGCAACACCTTCTCTGCTACTTCAGGGACCAAGTAT	488
QY	619	CCAGTTGTTGGAATGTCTCAACCCCTAGTATCTTCTGTTCCCAAGAGCTGTGCCCCC	678
Db	489	CCTCCCTAATGATGCTGTGCTGCTAGTCCCTAGTGCCTTCTGTTAGTACATCCTCAT	548
QY	679	CTGGCTAAAGGGGCTCCCTGTGTTATACAACTCTGCCTGCAATTTGCTCATCTCGAGCC	738
Db	549	GGAACTGGCCAGTCTCATTCAGCTTTTATCCATTCCTTATTTCTTCAACATTTGCCCT	608
QY	739	ACATTTGCCAAAGAGTTCTTCTTTTAGTAGATCTGGTCCAGGGTCACAACTTAAACACTAA	798
Db	609	GCATCATCTTTAGCCTGATGATGGAGGATTTGGTGTCTAGTATCCAGAGGGCCAG	668
QY	799	TTACAAAGGCACAGTCATTTGATGTGGCCAGTGTCCACAG-----	841
Db	669	TCTCTGATTGATTTAGGATCTAGTAGCTCAACTTCTCTCAACTGCTTCCCTCTCAGGGA	728
QY	842	-----TGGCAGAGTGGCTGTTCTTCAGTCATCAGACTGAAATACAGG	885
Db	729	TCACCTAAGACAGGACCTCAGAGTGGGCGAGTTCCCTCAGGCTTCAAGATTAAGTATCGG	788
QY	886	CAATATTCAATAGTCATGACAAACTATGAGTGGACACTTAAACAGTCCCGCCCAAGCAAG	945
Db	789	CAAAAATTTAATAGTCTAGCAAAAGCATGAGCGGATACCTCTCAGGTTTCAAGCTAGA	848
QY	946	ACTATTCTTATGCACTCAAGTTTACACAGGCTCAGCTGGCTTCAATATGGAATCTTTCT	1005
Db	849	AATGCCCTTCTTCACTCAAACTCTCTCTCAAACTCAGCTAGTACTATTTGGACTCTGGCT	908
QY	1006	GACATTCATCAGATGGAACACTTTACAGCAGAGAAATTTATCTCGGCAATGCACCTCAT	1065
Db	909	GACATCGATGGTGCAGACAGTGTGAAGCTGGAAGATTTTATCTTGGCGATGCACCTCACT	968
QY	1066	GATGTAGCTATGTCTGGCCAACTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT	1125
Db	969	GACATGGCCAAAGCTGGACACCCACTACCACTGAGTTGCTCCGAGCTGTGCTCCCA	1028
QY	1126	TCTTTTAGAGAGTTCGATCTGGCAGTGGTATATCTGTCTATAGCTCAACATCTGTAGT	1185
Db	1029	TCCTTCAGAGGGGAAAGCAAGTTGAT-----TCTGTAAATGGAACCTGCTCTCATAT	1082
QY	1186	CAGAGGCTACAGAGGACCACTTTTAGAGATGAACACAACTTAGAAAGAAATTA	1245

RESULT 11
US-09-907-969-72
; Sequence 72, Application US/09907969
; Publication No. US20030091580A1
; GENERAL INFORMATION:
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: King, Gordon E.
; APPLICANT: Algate, Paul A.
; APPLICANT: Fling, Steven P.

Db	1779	TTACTTCTATAAAGTCAAGAAAAGGAAGAAATATATGCGAAGAGCTTAAAGACAAATTA	1838
Qy	1966	GATGAAGTGGAGAAAGAACTAGATCAAAAGCTACAGGAGATTGATATTTTCAATAATCAG	2025
Db	1839	GATGCTCTTGAAGAAAGAACTGCATCTAAGCTCTCAGAAATGGAATTCATTTAAACAATCAG	1898
Qy	2026	CTGAAGGAATTAAGAAATACACATAAGCAACAACTCCAGAAAGCAAAAGTCCATGGAG	2085
Db	1899	CTGAAGGAATCTCAGAAAGAGCTATATAACACAGCAGTTAGCCCTTGAACAACATTCATANA	1958
Qy	2086	GCTGAACGACTGAACACAGAAAGAAACAAAGAACGAAAGATCATAGAAATTAGAAAAACAAA	2144
Db	1959	ATCAAAAGCTGACAAATTTGAAGGAATCGAAGAAAGAAAGATTAGACGCAAAAAA	2017
RESULT 12			
US-09-827-271-72			
; Sequence 72, Application US/09827271			
; Publication No. US20030165504A1			
; GENERAL INFORMATION:			
; APPLICANT: Retter, Marc W.			
; APPLICANT: Fanger, Gary R.			
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND			
; TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER			
; FILE REFERENCE: 210121.462C6			
; CURRENT APPLICATION NUMBER: US/09/827,271			
; CURRENT FILING DATE: 2001-04-04			
; NUMBER OF SEQ ID NOS: 461			
; SOFTWARE: FastSeq for Windows Version 3.0			
; SEQ ID NO 72			
; LENGTH: 2017			
; TYPE: DNA			
; ORGANISM: Homo sapien			
US-09-827-271-72			
Query Match 9.8%; Score 507.8; DB 3; Length 2017;			
Best Local Similarity 55.8%; Pred. No. 4.9e-107;			
Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5;			
Qy	175	GGCGTCGATTTAGCAAGGTAAAGTAAACAGAACCATGGCTCAGTTTCCAACACCTTTTGGT	234
Db	9	GGCTGAGAGCTGCAAGAAAGATCAGGATCATGATGCTCAGTTTCCACAGCGATGAAT	68
Qy	235	GGCAGCCTGGATATCTGGGCATAACTGTATAGAGAAAGAGGAGGATGATCAGCAGTTTC	294
Db	69	GGAGGCCAAATATGTGGGCTATTACATCTGAAGAACGTACTAAGCATGATAAACAGTTT	128
Qy	295	CATAGTTTAAAGCCAAATCTGGATTCATTCTGATGATCAAGCTCAAGAACTTTT	354
Db	129	GATAACTCAACCTTCAGGAGGTATACATACAGGTATCAAGCCCGTACTTTTTCCTA	188
Qy	355	CAATCTGGGTATACCTCAACCTGTTTATAGCACAGATATGGGCATCTAGCTGACATGAATAT	414
Db	189	CAGTCAGCTCTGCGGCCCGCTTTTATGCTGAAATATGGGCTTATCAGATCTGAACAAG	248
Qy	415	GATGGAAAGATGGATCAAGTGGAGTTTCCATAGCTATGAACTTATCAAACTGAAGCTA	474
Db	249	GATGGGAAGATGGACCAAGAGTTCTCTATAGCTATGAACATCATCAAGTTAAAGTTG	308
Qy	475	CAAGGATATCAGCTACCTCTGCACTTCCCTCTGATGAACACAGCAACC---AGTTGCT	531
Db	309	CAGGGCCAAACAGCTGCTGTAGTCTCTCCCTCTATCATGAACAAACCCCTATGTTCTCT	368
Qy	532	ATTTCTAGCGCACCAAGATTTGGTATGGAGGTATCGCCAGCATGCCACCGCTTACAGCT	591
Db	369	CACTAATCTCTGCTGCTTTTGGGATGGAGAGCATGCCAATCTGTCTCATTCACAGCA	428
Qy	592	GTTGCTCCAGTGCCTAATGGGATC-----CATTT	618
Db	429	TGCTCCAGTTGCACCTTATAGCAACACCTTTGTTCTGCTACTTTCAGGGACCAAGTAT	488
Qy	619	CCAGTTGTTGAATGTCTCAACCTAGTATCTTCTGTTCCACAGCAGCTGTGCCCCC	678

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QY 1726 AGCGAAGAAATTAGAGAGCAAAACAAATCTAGAGAGTTGAGAAATGCGGAAATCAACCAT 1785
Db      |||
1599 AAGACTGAGCTAGAAAGTTTGGATAAACAGTGTGACCTGGAAATATGGAATCAACAA 1658
QY      |||
1786 CTACAGCACAATTAACAGAACTCAGCAAAATGCTTGGAGACTTATCCAGAAACAG 1845
Db      |||
1659 CTTCAACAAGAGCTTAAGGAATATCAAAATAAGCTTATCTATCTGGTCCCTGAGAAGCAG 1718
QY      |||
1846 ATACTCAATGACCAATTAACAAAGTTTACAGCAACAGTTTGCACAGAGATTCACCTTGT 1905
Db      |||
1719 CTATTAAACGAAGAATTAACAACTAGCAGCTAGTAAACACCTGATTCAGGGATCAGT 1778
QY      |||
1906 ACACCTTAAAGAGCTTGAAGCAAAAGAACTAGCTCGGAGCACTACAGAGACCACTG 1965
Db      |||
1779 TTACTTTCATAAAAGTCATCAGAAAAGGAAGAAATATGCAAGACTTAAAGAAACAATTA 1838
QY      |||
1966 GATGAAGTCGAGAAAGAACTAGATCAAACTACAGAGATTCATATTTCAATATCAG 2025
Db      |||
1839 GATGCTCTTTGAAAAGAACTGCATCTAAGCTCTCAGAAATGGAATTCATTTAACAAATCAG 1898
QY      |||
2026 CTGAAGGAATAGAGAAATACAAATAAGCAACAACTCCAGAAAGCAAAAGTCCATGGAG 2085
Db      |||
1899 CTGAAGGAATCAGAGAAAGCTATATACACAGCAGTTAGCCCTTGAACAACTTCATAAA 1958
QY      |||
2086 GCTGAACGACTGAACAGAAAGAAACAAAGAACGAAAGATCATAGAAATAGAAAACAAA 2144
Db      |||
1959 ATCAAAAGTCGACAAATTTGAAGGAATCGAAAGAAAGAAATAGAGCAAAAGAAAAA 2017

RESULT 13
US-10-198-053-72
; Sequence 72, Application US/10198053
; Publication No. US20030124140A1
; GENERAL INFORMATION:
; APPLICANT: Bangur, Chaitanya S.
; APPLICANT: Retter, Marc W.
; APPLICANT: Fanger, Gary R.
; APPLICANT: Hill, Paul
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
; FILE REFERENCE: 210121.462C9
; CURRENT APPLICATION NUMBER: US/10/198,053
; CURRENT FILING DATE: 2002-07-17
; NUMBER OF SEQ ID NOS: 624
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 72
; LENGTH: 2017
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-198-053-72

Query Match 9.8%; Score 507.8; DB 6; Length 2017;
Best Local Similarity 55.8%; Pred. No. 4.9e-107;
Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5;

QY 175 GCGTCGATTTAGCAAGGTAAAGTAACAGAACCATCGCTCAGTTTCCACACCTTTTGGT 234
Db      |||
9 GCGTGAGAGCTGCAAGAAAGAGTCAGGATCATGATGGCTCAGTTTCCACAGCGATGAAT 68
QY 235 GCGAGCTCGATATCTGGGCCATAA CTGTAGAGAAAGAGCGAAGCATGATCAGCAGTTTC 294
Db      |||
69 GGAGGGCCCAATATGTGGGCTATTATCATCTGAAGAACGTACTAAGCATGATAAACAGTTT 128
QY 295 CATAGTTTAAAGCCAATATCTGATTCATTACTGTGTGATCAAGCTAGAAACTTTTTTTTT 354
Db      |||
129 GATAA CTTCAACCTTCAGGAGGTATACATAACAGGTGATCAAGCCGCTACTTTTTCCTA 188
QY 355 CAATCTGGGTACTCTCAACCTGTTTTAGCACAGATATGGGCATAGCTGACATGAATAT 414
Db      |||
189 CAGTCAGGTCTGCGGCCCCGGTTTTAGCTGAAATATGGGCCCTTATCAGATCTGAAACAAG 248
QY 415 GATGGAAGAATGGATCAAGTGGAGTTTTCATAGCTATGAAACTTATCAAACTGAAGCTA 474
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Db      |||
249 GATGGAAGATGGACCACCAAGAGTTCTCTATAGCTATGAACCTCATCAAGTTAAAGTTG 308
QY      |||
475 CAAGGATATCAGCTACCTCTGCACTTCCCTCTGTCATGAAACAGCAACC---AGTTGCT 531
Db      |||
309 CAGGCCCACACAGCTGCTGTAGTCTCTCTCTATCATGAACCAACCCCTATGTCTCT 368
QY      |||
532 ATTTCTAGCGCACCAAGCAATTTGGTATGGAGGTATCGCCAGCATGCCACCGCTTACAGCT 591
Db      |||
369 CCACCTAAATCTCTGCTGCTTTTGGGATGGGAAGCATGCCCAATCTGTCCATTCATCAGCCA 428
QY      |||
592 GTTGCTCTCAGTGCCTAATGGGATC-----CATTT 618
Db      |||
429 TTGCCCTCAGTGTGACCTATAGCAACACCTTGTCTTCTGCTACTTTCAGGGACCAGTATT 488
QY      |||
619 CCAGTTGTTGGAATGCTCCAAACCTAGTATCTTCTGTTCACAGAGCTGTGCCCCCCC 678
Db      |||
489 CTTCCCTTAATGATGCCTGCTCCCTAGTGCCTTCTGTGTAGTACATCTCTCATTTAGCAAT 548
QY      |||
679 CTGGGCTAAGGGGCTCCCTGTTTATACAACTCTGCCCTGCATTTGCTCATCTCCAGCC 738
Db      |||
549 GGAACCTGCCAGTCTCATTTACGCTTTATCCATTTCTTCTTCAACATTTGCTCTCAT 608
QY      |||
739 ACATTTGCCAAAGAGTTCTTCCCTTATAGTAGATCTGCTCCAGGGTCAACACTAAACACTAAA 798
Db      |||
609 GCATCATCTTACAGCTGATGATGGAGGATTTGGTGGTGTAGTATCCAGAGGCCCCAG 668
QY      |||
799 TTACAAAAGGCACAGTCAATTTGATGTGCCAGTGTGCCACACAG-----841
Db      |||
669 TCTCTGATTTAGTATTAGGATCTAGTAGCTCAACTCTCACTGCTTCCCTCTCAGGGAAC 728
QY      |||
842 -----TGGCAGAGTGGGCTGTTCTCTAGCTCATCAAGACTGAAATACAGG 885
Db      |||
729 TCACCTAAGACAGGAGCCTCAGAGTGGGCGAGTTCTCTCAGCCTTCAAGATTTAAGTATCGG 788
QY      |||
886 CAATTTATCAATAGTCATGACAAAACTATAGTGTGACACACTTAAACAGGTCCCAACAGCAAG 945
Db      |||
789 CAAAATTTTATAGTCTAGACAAAGGCATGAGCGATACCTCTCAGGTTTTCAGAGTAGA 848
QY      |||
946 ACTATTCTATGCAAGTTTACACAGGCTCAGGCTTCAATATGGAATCTTTCT 1005
Db      |||
849 AATGCCCTTCTCAGTCAAACTCTCTCAAACTCAGTAGCTACTATTTTGGACTCTGGCT 908
QY      |||
1006 GACATTTGATCAAGATGGAAACTTACACAGAGGAATTTATCTCGCAATGCACTCATTT 1065
Db      |||
909 GACATCGATGGTGACGACAGTTGAAAGCTGAAGATTTATTCTGCGGATGACCTCATCT 968
QY      |||
1066 GATGTAGCTATGCTGGCCCAACCACTGCCACCTGTCTGCTCTCCAGAAATACATTCACCT 1125
Db      |||
969 GACATGGCCAAAGCTGGACAGCCACTACCCTGACGTTGCTCTCCGAGCTTGTCCCTCCA 1028
QY      |||
1126 TCTTTTGAAGAGTTTCGATCTGGCAGTGTATATCTGTATAGCTCAACATCTGTAGAT 1185
Db      |||
1029 TCTTTTCAAGGGGGGAAAGCAAGTTGAT-----TCTGTTAATGGAACCTCTGCTCTTCATAT 1082
QY      |||
1186 CAGAGCTTACCAGAGGAACCAAGTTTATAGAGATGAACAACAATTAGAAAAGAAATTA 1245
Db      |||
1083 CAGAAAACACAAGAGAAGAGAGCCT-----CAGAAGAACTG 1118
QY      |||
1246 CCGTTAAACGTTTGAAGATAAGAGGGGAGAACTTTTGAACGTGGCAACCTCGGAACCTGGAG 1305
Db      |||
1119 CCAGTTACTTTTGAAGCAACAAAGCCAACTATGAACGAGGAAACATGAGCTGGAG 1178
QY      |||
1306 AAACGAAGCAAGCTCTCTCTGGAACAGCAGCCCAAGGAGCAGGAGCGCTTGGCCCGCTG 1365
Db      |||
1179 AAGCGACGCAAGTGTGTATGGAGCAGCAGAGAGGGAGGCTGAAACGCAAAAGCCCAAGAA 1238
QY      |||
1366 GAGCGGCGGACAGAGAGGAGAGCGGTAGCGCCAGAGCAAGAGCGCAAAAGACAA 1425
Db      |||
1239 GAGAAGGAAGTGGAGCGGAAACAGAGAACTTGCAGAGCAAGATGGGAAGAGCAG 1298
QY      |||
1426 CTGGAACCTGGAGAAGCACTGGAAAAGCAGCGGAGCTAGAACGCGAGAGAGAGGAGGAG 1485
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Db 1299 CTGGAGTTGGAGAAACGCTTGGAGAAAACAGAGAGAGCTGGAGACAGCGGAGGAGAG 1358
Qy 1486 AGGAGGAAAGAAATTGAGAGGCGAGAGGCTGCAAAAACGGGAACTTGAAGGCAACGACAA 1545
Db 1359 AGGAGAAAGGAGATAGAAAGACGAGAGGCGCAAAACAGGAGCTTGAGAGACAAACGCCGT 1418
Qy 1546 CTTGAGTGGGACCGGATCGAAGGCAAGGACACTACTAATCAAGAAACAAACAAACAGAG 1605
Db 1419 TTAGAATGGGAAAGACTCCGTCGGCAGAGCTGCTCAGTCAGAGACGAGGAAACAGAA 1478
Qy 1606 GACATAGTTGTACTGAAAGCAAAAGAAAGAACTTTGGAAATTTGAATTTAGAAGCTCTTAAT 1665
Db 1479 GACATTTGTCAGGCTGAGCTCCAGAAAGAAAGTCTCCACCTGGAAGCTGGAAGCAAT 1538
Qy 1666 GATAAAAGCATCAACTGAAAGGAACTTTCAAGATATCAGATGTCGATTGACCAACCA 1725
Db 1539 GGAACAACATCAGCAGATCTCAGGACAGCTACAAGATGTCGAATCAGAAAGCAACACAA 1598
Qy 1726 AGCAAGAAATTCAGAGCACAACAAATCTAGAGAGTTGAGAAATGCGCCGAATCAACCAAT 1785
Db 1599 AAGACTGAGCTAGAAGTTTGGATAAACACAGTGTGACCTGGAATTAAGAAATCAACAA 1658
Qy 1786 CTACACCAAAATTCAGGAATCTCAGCAAACTCTTGAAGACTTATTCAGAGAAACACAG 1845
Db 1659 CTTCAACAGAGCTTAAGGATATCAAAATAGCTTATCTATCTGCTCCCTGAGAGCAG 1718
Qy 1846 ATACTCAATGACCAATTAACAAAGTTTCAGAGCAACAGTTTGCACAGAGATTCACATTGTT 1905
Db 1719 CTATTAAACGMAAGAAATTAACAAACATGCAGCTCAGTAAACACACCTGATTCAGGGATCAGT 1778
Qy 1906 ACATTTAAAGAGCTTAGAAGCAAAAGAACTAGCTCGCAGCAGCACTACGAGACCAACTG 1965
Db 1779 TTACTTCATAAAAGTCATCAGAAAGGAAATTAATGCAAGAGCTTAAAGAACAAATTA 1838
Qy 1966 GATGAAGTGAGAGAAAGAACTAGATCAAACTACAGGAGATGATATTTTCAATAATCAG 2025
Db 1839 GATGCTCTTGAAAGAAAGAACTGATCTAAGCTCTCAGAAATGATTCATTTAAACATCAG 1898
Qy 2026 CTGAAGGAACTAGAGAAATPACAAATAAGCAACAACTCCAGAGCAAAAGTCCATGGAG 2085
Db 1899 CTGAAGGAACTCAGAGAAAGCTAATAACACAGCAGTTAGCCCTTGCAACAACTTCATAAA 1958
Qy 2086 GCTGAAGCACTGAAACAGAAAGCAAGACAGCAAGAAAGTATCAGAAATTTAGAAACAAAA 2144
Db 1959 ATCAACGTCGACAAATTTGAAGGAAATCGAAAGAAAGAAATTTAGAGCAAAAGAAAAA 2017

RESULT 14
US-10-860-790-72
; Sequence 72, Application US/10860790
; Publication No. US20050031634A1
; GENERAL INFORMATION:
; APPLICANT: Bangur, Chaitanya S.
; APPLICANT: Retter, Marc W.
; APPLICANT: Fanger, Gary R.
; APPLICANT: Hill, Paul
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
; TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
; FILE REFERENCE: 210121.462C11
; CURRENT APPLICATION NUMBER: US/10/860,790
; CURRENT FILING DATE: 2004-06-02
; NUMBER OF SEQ ID NOS: 624
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 72
; LENGTH: 2017
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-860-790-72

Query Match 9.8%; Score 507.8; DB 8; Length 2017;
Best Local Similarity 55.8%; Pred. No. 4.9e-107;
Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5;

Qy 175 GCGCTCGATTAGCAAGGTAAGAAAGTAAACAGAAACCATGGCTCAGTTTCCCAACACACTTTTGGT 234
Db 9 GGTGAGAGCTGCAAGAGAGAGTCAAGATCATGATGGCTCAGTTTCCCAACAGCATGAAT 68
Qy 235 GGCAGCTGGATATCTGGGCCATTAATCTGTAGAGGAAAGAGCGAAGCATGATCAGCAGTTTC 294
Db 69 GGAGGCCAAATATCTGGGCTATTACATCTGAAGAACGCTACTAAGCATGATATAACAGTTT 128
Qy 295 CATAGTTTAAAGCCAAATATCTGGATTCAATCTGCTGATCAAGCTAGAGAACTTTTTTTTTT 354
Db 129 GATAACCTCAAACTTCAAGAGGTACATACAGGTGATCAAGCCGCTATTTTTCCTTA 188
Qy 355 CAATCTGGGTTACCTCAACCTGTTTACACAGATATGGCAGTCTAGCTGACATGAATAT 414
Db 189 CAGTCAGGCTGCGCGGCCCGGTTTTAGCTGAAATATGGCCCTTATCAGATCTGAACAAG 248
Qy 415 GATGAAGATGATCAAGTGGAGTTTCCATAGCTATGAAACTTATCAAACTGAAGCTA 474
Db 249 GATGGGAAGATGGACAGCAAGAGTTCTCTATAGCTATGAAACTCATCAAGTTAAAGTTG 308
Qy 475 CAAGGATATCAGCTACCTCTGCACTTCCCTCTGTCATGAAACAGCAACC---AGTTGCT 531
Db 309 CAGGGCAACAGCTGCTGCTGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 368
Qy 532 ATTTCTAGCGCACCAAGCAATTTGGTATGGAGGATATGGCAGATGCAACCGCTTACAGCT 591
Db 369 CCACATACTCTGCTCGTTTTGGGATGGGAGCATGCCCAATCTGTCCATTCATCAGCCA 428
Qy 592 GTTGCTCCAGTGGCAATGGGATC-----CAAT 618
Db 429 TTGCTCTCCAGTTGCACTATAGCAACACCTTTGCTCTTCTGCTACTTTCAGGGACGATAT 488
Qy 619 CCAGTTGTTGGAATGCTCCAACTTAGTATCTCTGTTCCCAACAGCAGCTGTGCCCTCC 678
Db 489 CTTCCCTAATGATGCTGCTCTCCCTAGTGGCTTCTGTTAGTACATCTCTCTCTCTCT 548
Qy 679 CTGGCTAAACGGGCTCTCCCTCTGTTATACAACTCTGCTGCAATTTGCTCATCTGAGCC 738
Db 549 GGAACCTGCCAGTCTCATTCAGCTTTATCCATTCCTTATCTCTTCAACATTTGCTCAT 608
Qy 739 ACATGCGCAAGAGTTTCTCTTTTAGTATCTGCTCAGGCTCAGGTCACAACTAACACTAA 798
Db 609 GCATCATCTTACAGCTGATGATGGAGGATTTGGTGGTGTAGTATCCAGAAAGGCCAG 668
Qy 799 TTACAAAGGCACAGTCATTTGATGCGCAGTGTCCCAACCAG----- 841
Db 669 TCTCTGATTTAGGATCTAGTAGCTCAATCTCTCACTGCTTCCCTCTCAGGGAAC 728
Qy 842 -----TGGCAGAGTGGGCTGTTCTCAGTCATCAAGACTGAAATACAGG 885
Db 729 TCACCTTAAGACAGGGACCTCAGAGTGGGAGTTCTCTCAGCTTCAAGATTTAAAGTATCG 788
Qy 886 CAATTAATCAATAGTATGACAAACTATGATGGACACTTAACAGGTCCCCAAGCAAGA 945
Db 789 CAAAAATTTAATAGTCTAGACAAAGGCATGAGCGGATACCTCTCAGGTTTTCAGCTAGA 848
Qy 946 ACTATTTCTTATGCACTCAAGTTTACCACAGGCTCAGCTGGCTTCAATATGGAATCTTTCT 1005
Db 849 AATGCCCTTCTTCAAGTCAAAATCTCTCTCAACTCAGTAGTACTATTTGGACTCTGGCT 908
Qy 1006 GACATTTGATCAAGATGGAAGAACTTACAGCAGAGGAAATTTATCTGGCAATGCACTCAT 1065
Db 909 GACATCGATGGTGCAGGACAGTTGAAAGCTGAAAGATTTATCTGGGATGCACTCACT 968
Qy 1066 GATGTAGCTATGCTGGCCAAACCACTGCCACCTGCTGCTGCTCAGATACATTTCCACCT 1125
Db 969 GACATGGCCAAAGCTGGACAGCCACTACCACTGACGTTGCTTCCGAGCTTGTCCCTCCA 1028
Qy 1126 TCTTTTAGAGAGTTCGATCTGCGAGTGGTATATCTGTCTATAAGCTCAACATCTGTAGAT 1185
Db 1029 TCTTTAGAGGGGAAAGCAAGTTGAT-----TCTGTTAATGNACTCTGCTCTCATAT 1082
Qy 1186 CAGAGGCTACCAGAGGAACCAAGTTTATAGAGATGAACAAACAAATTTAGAAAAGAAATTA 1245


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Db 1083 CAGAAAAACAAGAAAGAGCGCT-----CAGAAAGAACTG 1118
Qy 1246 CTGTGAACCTTTGAAGTAAGACGGGAGAACTTTGAACGTGGCAACCTGGAACCTGGAG 1305
Db 1119 CCAGTTACTTTTGAAGACAAACGGAAGCCACTATGAACGAGGAAACATGAGCTGGAG 1178
Qy 1306 AAACGAGGCAAGCTCTCTCTGGAACAGACGCGCAAGGACGAGGAGCGCTGGCCCCAGCTG 1365
Db 1179 AAGCGACGCAAGTGTGTGATGAGCAGCAGCAGAGGAGGCTGAACGCCAAGGCCAGAAA 1238
Qy 1366 GAGCGGCGGAGCAGAGAGGAGGAGCTGTGAGCGCCAGGACGAGAGCGCAAAAGACAA 1425
Db 1239 GAGAAGGAAGTGGGAGCGGAAACAGAGAGAACTGCNAAGACGCAAGATGGAAGAGACG 1298
Qy 1426 CTGGAACTCGAGAAGCAACTGGAAAAGCAGCGGAGCTAGAACGGCAGAGAGAGGAGAG 1485
Db 1299 CTGGAGTTGAGANAACGCTTGGNAAACAGAGAGAGCTGGAGAGCAGCGGAGGAGAG 1358
Qy 1486 AGGAGGAAAGAAATTGAGAGGCGAGAGCTTGCAAAAACGGGAACCTTGAAGGCAACGACAA 1545
Db 1359 AGGAGAAAGGAGATAGAAAGACGAGAGGCGCAAAACAGGAGCTTGAGAGACAAACGCCGT 1418
Qy 1546 CTTGAGTGGGAACGGAATCGAAGGCAAGAACTACTTAATCAAGAAACAAAGACAGAG 1605
Db 1419 TTGAATGGGAAGACTCCGTCGGCAGGAGCTGCTCAGTCAGAAGACCGAGGAAACAGAA 1478
Qy 1606 GACATAGTTGTACTGAAAACGAAGAAAAGACTTTTGGAAATTTGAAATTTAGAACTCTAAAT 1665
Db 1479 GACATTTGTCAGGCTGAGCTCCAGAAAGAAAGTCTCCACTGGAAGCTGAGAGAGTGAAT 1538
Qy 1666 GATAAAAGCATCAACTAGAGGGAACCTTCAAGATATCAGATGTGCGATTTGACCCCAAA 1725
Db 1539 CGAAACATCAGCAGATCTCAGCAGACTACAAGATGTCCAAATCAGAAAGCAACACAA 1598
Qy 1726 AGGCAAGAAATTGAGAGCACAACAAATCTAGAGAGTTGAGAAATTCGCAAAATCAACAT 1785
Db 1599 AAGACTGAGCTAGAAAGTTTGGATAAACAGTGTGACCTGGAAATTTATGGAATTCAAACAA 1658
Qy 1786 CTACAGCAACAATTACAGGAATCTCAGCAAAATGCTTGGAAAGACTTTATCCAGAAAACAG 1845
Db 1659 CTTCAACAGAGCTTAAGGAATATCAAAATAGCTTATCTAATCTGCTCCTGAGAGCAG 1718
Qy 1846 ATACTCAATGACCAATTAAAAACAAGTTTACAGCAACAGTTTGCACAGAGATTCATGTT 1905
Db 1719 CTATTAAACGAAAGAAATTAATAACATGCAGCTCAGTAACACACCTGATTCAGGGATCACT 1778
Qy 1906 ACATTTAAAGAGCCTTAGAAGCAAAAGAACTAGCTCGCAGCAGCCTACAGACCAACTG 1965
Db 1779 TTACTTTCATAAAAGTCAATCAGAAAAGGAAGAAATTTATGCCAAGACTTAAAGAACAAATTA 1838
Qy 1966 GATGAAGTGGAGAAAGAACTAGATCAAAACTACAGGAGATTTGATATTTTCAATATACAG 2025
Db 1839 GATGCTCTTGAAGAAAGAACTGCATCTAAGCTCTCAGAAATGGATTCATTTAACAATCAG 1898
Qy 2026 CTGAAGGAACCTAAGAAATACACAATAAGCAACAACTCCAGAGCAAGAAAGTCCATGGAG 2085
Db 1899 CTGAAGGAACCTCAGAGAAAGCTATATAACACAGCAGTTAGCCCTTGAACAACTTCATAAA 1958
Qy 2086 GCTGACGACTGAAACAGAAAGCAAGCAAGCAAGAGATCATAGAAATTTAGAAAACAAA 2144
Db 1959 ATCAAAACGTGACAAATTGAAGGAATTCGAAAGAAAAGAAATTTAGAGCAAAAACAAAAA 2017
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RESULT 15

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US-09-764-881-55
; Sequence 55, Application US/09764881
; Publication No. US20020086821A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
; FILE REFERENCE: PT207
; CURRENT APPLICATION NUMBER: US/09/764,881
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; CURRENT FILING DATE: 2001-01-17
; Prior application data removed - refer to PALM or file wrapper
; NUMBER OF SEQ ID NOS: 192
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 55
; LENGTH: 568
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (481)
; OTHER INFORMATION: n equals a,t,g, or c
; NAME/KEY: SITE
; LOCATION: (536)
; OTHER INFORMATION: n equals a,t,g, or c
; NAME/KEY: SITE
; LOCATION: (556)
; OTHER INFORMATION: n equals a,t,g, or c
; NAME/KEY: SITE
; LOCATION: (562)
; OTHER INFORMATION: n equals a,t,g, or c
US-09-764-881-55
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Query Match          9.7%; Score 503.6; DB 3; Length 568;
Best Local Similarity 96.9%; Pred. No. 2.1e-106;
Matches 554; Conservative 0; Mismatches 13; Indels 5; Gaps 4;

Qy 130 ACTGATTTGTCCCTGGGCGGCGAGCGCGGACCCCGCGAGATGAGGCGTCGATTAGCAA 189
Db 1 ACTGATTTGTCCCTGGGCGGCA-CGCGGACCCCGCGAGATGAGGCGTCGATTAGCAA 59

Qy 190 GGTAAAAAGTAACAGAACCATGCTCAGTTTCCAAACCTTTTGGTGGCAGCCTGGATATC 249
Db 60 GGTAAAGTAACAGAACCATGCTCAGTTTCCAAACCTTTTGGTGGCAGCCTGGATATC 119

Qy 250 TGGGCCATAACTGTAGAGAAAGCAGAGCATGATCAGCAGTTCCATAGTTTAAAGCCA 309
Db 120 TGGGCCATAACTGTAGAGAAAGCAGCA--CATGATCAGCAGTTCCATAGTTTAAAGCCA 177

Qy 310 ATATCTGGATTCATTACTGCTGATCAAGCTAGAAACCTTTTTTTTCAATCTGGGTACCT 369
Db 178 ATATCTGGATTCATTACTGCTGATCAAGCTAGAAACCTTTTTTTTCAATCTGGGTACCT 237

Qy 370 CAACCTGTTTTAGCAGATATGGGCACCTAGCTGACATGAATATATGATGGAAGATGGAT 429
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Qy 430 CAAAGTGGAGTTTTCCCATAGCTATGAAACTTATCAAACTGAAGCTACAAGGATATCAGCTA 489
Db 298 CAAAGTGGAGTTTTCCCATAGCTATGAAACTTATCAAACTGAAGCTACAAGGATATCAGCTA 357

Qy 490 CCCTCTGCACITCCCCCTGTTCATGAACAGCAACCAAGTTGCTATTCTAGCGCACAGCA 549
Db 358 CCCTCTGCACITCCCCCTGTTCATGAACAGCAACCAAGTTGCTATTCTTAGCGCACAGCA 417

Qy 550 TTTGTGATGGAGAGTATCGCCAGCATGCCCGCTTACAGCTGTTGCTCCAGTGCCAATG 609
Db 418 TTGTGATGGGAGGTATCGCCAGCAAGCCCGCTTACAGCTGTTGCTCCAGTGCCAATG 477

Qy 610 GGTATCCATTTCCAGTTGTT--GGAATGCTTCCAAACCTTAGTATCTTCTGTTCCACAGCAGC 668
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Qy 669 TGTGCCCCCCTGCTGCTAAACGGGGCTCCCCCTG 700
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GenCore version 5.1.7
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OM nucleic - nucleic search, using sw model

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Gapop 10.0 , Gapext 1.0

Searched: 1303057 seqs, 888780828 residues

Total number of hits satisfying chosen parameters: 2606114

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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9: /cgn2_6/ptodata/1/ina/backfileseq.*

pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	ID	Description
1	551.6	10.6	5813	3	US-09-949-016-1739
2	507.8	9.8	2017	3	US-09-404-879A-72
3	507.8	9.8	2017	3	US-09-338-933-72
4	507.8	9.8	2017	3	US-09-215-681-72
5	507.8	9.8	2017	3	US-09-216-003A-72
6	507.8	9.8	2017	3	US-09-667-857-72
7	507.8	9.8	2017	3	US-10-198-053-72
8	507.8	9.8	2017	3	US-09-827-271-72
9	174	3.3	174	3	US-09-513-999C-27927
10	165.2	3.2	2873	3	US-08-630-915A-193
11	165.2	3.2	2873	3	US-09-879-957-193
12	163.8	3.2	747	3	US-08-630-915A-39
13	163.8	3.2	747	3	US-09-879-957-39
14	153.4	3.0	531	3	US-09-404-879A-5
15	153.4	3.0	531	3	US-09-338-933-5
16	153.4	3.0	531	3	US-09-215-681-5
17	153.4	3.0	531	3	US-09-216-003A-5
18	153.4	3.0	531	3	US-09-667-857-5
19	153.4	3.0	531	3	US-10-198-053-5
20	153.4	3.0	531	3	US-09-827-271-5
21	149.2	2.9	480	3	US-09-404-879A-60
22	149.2	2.9	480	3	US-09-338-933-60
23	149.2	2.9	480	3	US-09-215-681-60
24	149.2	2.9	480	3	US-09-216-003A-60

25	149.2	2.9	480	3	US-09-667-857-60	Sequence 60, Appl
26	149.2	2.9	480	3	US-10-198-053-60	Sequence 60, Appl
27	149.2	2.9	480	3	US-09-827-271-60	Sequence 60, Appl
28	140.6	2.7	531	3	US-09-404-879A-4	Sequence 4, Appl
29	140.6	2.7	531	3	US-09-338-933-4	Sequence 4, Appl
30	140.6	2.7	531	3	US-09-215-681-4	Sequence 4, Appl
31	140.6	2.7	531	3	US-09-216-003A-4	Sequence 4, Appl
32	140.6	2.7	531	3	US-09-667-857-4	Sequence 4, Appl
33	140.6	2.7	531	3	US-10-198-053-4	Sequence 4, Appl
34	140.6	2.7	531	3	US-09-827-271-4	Sequence 4, Appl
35	117	2.3	129127	3	US-09-949-016-13481	Sequence 13481, A
36	112.6	2.2	7218	2	US-08-232-463-14	Sequence 14, Appl
37	97	1.9	381	3	US-09-404-879A-61	Sequence 61, Appl
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41	97	1.9	381	3	US-09-667-857-61	Sequence 61, Appl
42	97	1.9	381	3	US-10-198-053-61	Sequence 61, Appl
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44	83	1.6	612	3	US-09-902-540-1357	Sequence 1357, Ap
45	82.6	1.6	1827	3	US-09-270-767-1308	Sequence 1308, Ap

ALIGNMENTS

RESULT 1
US-09-949-016-1739
; Sequence 1739, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CL001307
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 1739
; LENGTH: 5813
; TYPE: DNA
; ORGANISM: Human
US-09-949-016-1739

Query Match	10.6%	Score 551.6;	DB 3;	Length 5813;
Best Local Similarity	50.8%;	Pred. No. 4.4e-121;		
Matches 1812;	Conservative	0;	Mismatches 1594;	Indels 164; Gaps 14;
Qy	184	TAGCAAGTAAAGTAAACAGAACATGGCTCAGTTTCCAAACACCTTTGGTGGCAGCTG	243	
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Qy	244	GATATCTGGGCCATAACTGTAGAGGAAGAGCAAGCATGATCAGCAGTTCCATAGTTTA	303	
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Qy	304	AAGCAATATCTGGATTCATTACTGGTGATCAAGCTAGAACTTTTTTTTCAATCTGGG	363	
Db	138	AAACCTTCAGGAGGTTTACATAACAGGTGTATCAAGCACGTATTTTCTCAATCAGGT	197	
Qy	364	TTACTCTCAACTGTTTTAGCACAGATATGGCCTAGCTAGCATGAATAATGATGGAAGA	423	
Db	198	CTGGCGCCCTGTTTTAGCTGTAATATGGCTTTATCAGACCTAAACAGGATGGGAAG	257	
Qy	424	ATGGATCAAGTGGAGTTTTCATAGCTATGAAACTTATCAAACTTGAAGCTCAAGGATAT	483	

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Db	318	CAGTTGCTGTGGTTCTCCCTCTCTATATGAAGCAACCCCTATGTTTCTCATTAAT	377	
Qy	541	GCACGACATTTGGTATGGAGGTATCGCCAGCATGCCCGGTACAGGTGTGTGCTCA	600	
Db	378	TCTGCTCGTTTGGAAATGGGAAGCATGCCCAATCTGTCCATTCCTCAGCCATTTGCCCTCA	437	
Qy	601	GTGCCAATGGGATC-----	CAATCCAGTTGTGA	630
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Qy	631	ATGTTCTCCAACCTTAGTATCTTCTGTCCACAGCAGCTGTGCCCCCTCGGTAAACGGG	690	
Db	498	ATGCCACCTCCCTAGTGCTTCTGTAGCACATCATATTACCAATGGAACCGCCAGT	557	
Qy	691	GCTCCCTCTGTTATACAACTCTGCCCTGCAATTTGCTCATCTCGCAGCCACATTTGCCAAG	750	
Db	558	CTCAITCAGCCCTTTACCCATTCCTTATCTTCTCAACATTTGCCCTCATGCTTAT	617	
Qy	751	AGTCTCTCTTAGTAGATCTGTCACAGGTCACTAACTAACTAAATTTACAAAGGCA	810	
Db	618	AGTCTGATGATGGAGGATTTGGAGGTGTAGTATACAGAAAGCGAGTCTCTGATTGAT	677	
Qy	811	CAGTCAATTTGATGTGCCAGTGTCCACCAG-----	841	
Db	678	TTAGGATCTAGTAGCTCAACTCTCTCGACTGCTTCACTCTCAGGGAACCTCACCAAGACT	737	
Qy	842	----TGGCAGATGGCTGTCTCTAGTCAATCAAGACTGAATACAGGCAATTTTCAT	897	
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Qy	1558	CGGAATCGAAGCGCAAGAACTACTAAATCAAAAGAAACAAAGCAAGAGGACATAGTTGA	1617	
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Qy	1918	GCCTTAGAAGCAAAAGAACTAGCTCGGAGCACTTACGAGACCACTTGGATCAAGTGGAG	1977	
Db	1788	AAATCATTTAGAAAGGAAGAAATTTGCCAAAGACTTTAAGAACAGTTAGATGCTCTTGAA	1847	
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Db	1848	AAAGAACTGCATCTAAGCTGTCAAGATGGATTTCTTTTAACTCACTTAAAGGAACCTG	1907	
Qy	2038	AGAGAAATACACAATAAGCAACACTCCAGAAAGAAAGTCCATGGAGGCTGAAACGACTG	2097	
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Qy	2098	AAACAGAAAGCAAGAAAGCAAGAACTATAGAAATTT-----AGAAAAACAAAAAGAA	2148	
Db	1968	AAATTTGAGGAATTTGAAAGGAAGAAATTTAGAACTTATGCAGAAAAAGAACTAGAGAT	2027	
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QY 1546 CTGTAGTGGCAACGGAATCGAAGGCAAGAACTACTAAATCAAAGAAACAAAGCAAGAG 1605
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QY 2086 GCTGAACGACTGAACAGAGAAAGCAAGAAAGATCATAGAATTTAGAAAACAAA 2144
Db 1959 ATCAAACTGACAAATTTGAAGGAATCGAAAGAAAAGATTAGAGCAAAAAA 2017
RESULT 3
US-09-338-933-72
; Sequence 72, Application US/09338933
; Patent No. 6488931
; GENERAL INFORMATION:
; APPLICANT: Mitcham, Jennifer Lynn
; APPLICANT: King, Gordon E.
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY OF
; TITLE OF INVENTION: OVARIAN CANCER
; FILE REFERENCE: 210121.462C1
; CURRENT APPLICATION NUMBER: US/09/338,933
; CURRENT FILING DATE: 1999-06-23
; NUMBER OF SEQ ID NOS: 312
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 72
; LENGTH: 2017
; TYPE: DNA
; ORGANISM: Homo sapien
US-09-338-933-72
Query Match 9.8%; Score 507.8; DB 3; Length 2017;
Best Local Similarity 55.8%; Pred. No. 7.3e-111;
Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5;
QY 175 GCGCTGCTGATAGCAAGGTAAAGTAA CAGAACCATGGCTCAGTTTCCACACCTTTTGGT 234
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QY 295 CATAGTTTAAAGCCAAATATCTGGATTCATTACTGTGTGATCAAGCTAGAAACTTTTTTTT 354
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QY 475 CAAGGATATCAGTACCTCTGCACTTCCCTCTGTGATGAAACAGCAACCC---AGTTGCT 531
Db 309 CAGGCCCAACAGCTGCGCTGTAGTCCTCCTCTCATCATGAAACAAACCCCTATGTCTCT 368
QY 532 ATTTCTAGCGCACAGCATTTTGGTATGGAGGTATCGCCAGCATGCCACCGCTTACAGCT 591
Db 369 CCACTAATCTCTGCTGTTTGGGATGGGAAGCATGCCCAAATCTGTCCATTCATCAGCCA 428
QY 592 GTTGCTCCAGTGCCTCAATGGGATC-----CAATT 618

Db 429 TCGCTCCAGTGTGCACCTATAGCAACCCCTTGTCTTCTGCTACTTTCAGGACCAGTAT 488
Qy 619 CAGATTGTTGGATGCTCCAAACCTTAGTATCTTCTGTTCCACAGACAGCTGCCCCC 678
Db 489 CCTCCCTAATGATGCTGCTCCCTTAGTCTTCTGTTAGTACATCTCAATACCAAT 548
Qy 679 CTGGCTAAAGGGGCTCCCTGTTATACAACTCTGCTGCTATTTGCTCATCTGAGCC 738
Db 549 GGAACCTGCCAGTCTCATTCAGCCCTTTATCCATCTCTTATCTCTTCAACATTCGCTCAT 608
Qy 739 ACATTGCCAAAGAGTTCTTCTTTAGTATGATCTGCTCCAGGGTCACAACTAAACACTAAA 798
Db 609 GCATCATCTTAAGCTGATGATGGAGGATTTGGTGGTCTAGTATCCAGAGGCCAG 668
Qy 799 TTAACAAAGCAGCATCTTTGATGTGGCAGTGTCCCACAG-----841
Db 669 TCTCTGATTGATTAGGATCTAGTAGCTCAACTTCTCAACTGCTTCCCTCTCAGGGAAC 728
Qy 842 -----TGGCAGAGTGGCTGTTCTCAGTCTCATCAAGCTGGAATACAGG 885
Db 729 TCACCTAAGACAGGGACCTCAGAGTGGCGAGTTCTCAGCCTTCAAGATTTAAAGTATCGG 788
Qy 886 CAATTATTCATAGTATGACAAAACCTATGAGTGGACACTTAACAGCTGCCCAAGCAAGA 945
Db 789 CAAAAATTTAATAGTCTAGCAAGGATGAGCGGATACCTCTCAGGTTTCAAGCTAGA 848
Qy 946 ACTATTCTTATGAGTCAAGTTTACCAAGGCTCAGCTGGCTTCAATATGGAATCTTTCT 1005
Db 849 AATGCCCTTCTTCAAGTCAATCTCTCAAACTCAGCTAGCTACTATTTGGACTCTGGCT 908
Qy 1006 GACATTGATCAAGATGGAACCTTACAGAGAGGAATTTCTTGGCAATGCACTCAT 1065
Db 909 GACATCATGCTGACGACAGTGTGAAGCTGAAGATTTATCTGGCGATGCACTCACT 968
Qy 1066 GATGTAGTATGCTGCGCAACCACTGCCACCTGCTCTGCTCCAGAAATACATTCACCT 1125
Db 969 GACATGCCAAAGCTGGACAGCAGCTACCACTGAGTTGCTCCGAGCTGTGCCCTCA 1028
Qy 1126 TCTTTTGAAGAGTTCGATCTGGCAGTGGTATATCTGTCTAATAGCTCAACATCTGTAGAT 1185
Db 1029 TCTTTGAGGGGGAAGCAAGTTGAT-----TCTGTTAATGAACTCTGCTCTCATAT 1082
Qy 1186 CAGAGGCTACAGAGGAACCGTTTGAAGATGAACAACTATTAGAAAGNATTA 1245
Db 1083 CAGAAAAACAAGAAAGAGGCT-----CAGAAGAACTG 1118
Qy 1246 CCTGTAAGTTTGAAGATAGAGCGGAGAACTTTGAACGTGGCAACCTGGAACTGGAG 1305
Db 1119 CCAGTTACTTTTGGAGCAACCGAAAGCCACTATGAACAGGAACATGGAGCTGGAG 1178
Qy 1306 AAACGAAGCAAGCTCTCTCGAAACAGCAGCGCAGGAGGAGCGCTGGCCCACTG 1365
Db 1179 AAGCAGCCCAAGTTGATGGAGCAGCAGCAGGAGGAGCTGAACGCAAGCCAGAAA 1238
Qy 1366 GAGCGGCGGAGCAGAGAGAGAGAGCGGTGAGCGCAGGAGCAGAGCGCAAGAAAGCAA 1425
Db 1239 GAGAAGGAAGAGTGGGAGCGGAAACAGAGAGAACTGCAAGAGCAAGAAATGGAAGAGCAG 1298
Qy 1426 CTGGAACTGGAGAGCAACTGGAAAGCAGCGGAGCTAGAAACGCGAGAGAGAGGAGGAG 1485
Db 1299 CTGAGTTGGAGAAACGCTTGGAGAAACAGAGAGCTGGAGAGCAGCGGAGAGAGAG 1358
Qy 1486 AGGAGGAAGAAATTTGAGAGCGCAGAGGCTCAAAAACGGGAACTTGAAGGCAACGACAA 1545
Db 1359 AGGAGAAAGAGATAGAAAGACAGAGGAGGAGCAAAACAGGAGCTTGAAGAGCAACGCGT 1418
Qy 1546 CTTGAGTGGGAACGGAATGGAAGGAGAGAACTAATAATCAAGAAACAAAGAGACAGAG 1605
Db 1419 TTAGAAAGGAAAGAACTCCGTGGCAGGAGCTGCTCAGTCAGAGAGCCAGGGAAACAGAA 1478
Qy 1606 GACATAGTTGATGGAAGCAAGAAAGAACTTTGGAAATTTGAATTAGAGCTCTAAAT 1665

Db 1479 GACATTGTGAGCTGAGCTCCAGAAAGAAAGTCTCCACCTGGAACTGGAAGCAGTGAAT 1538
Qy 1666 GATAAAAGCATCAACTAGAACGGAAACTTTCAAGATATCAGATGTGATTTGACCCACCAA 1725
Db 1539 GAAAGACATCAGCAGATCTCAGCAGACTTCAAGATGTCCAAATCAGAAAGCAACACAA 1598
Qy 1726 AGGCAAGAAATTTGAGAGCACAACAAATCTAGAGAGTTGAGAAATTCGCCAAATCACCCAT 1785
Db 1599 AGACTGAGCTAGAAAGTTTGGATAAACAGTGTGACCTGGAAATTTATGGAAATCAACAA 1658
Qy 1786 CTACAGCAACAATTCAGGAATCTCAGCAAAATCTTGGAAAGACTTATTCAGAAAAACAG 1845
Db 1659 CTTCAACAGAGCTTAAGGAATATCAAAATTAAGCTTATCTATCTGCTCCTGGAAGCAG 1718
Qy 1846 ATACTCAATGACCAATTTAAACAAAGTTTCAAGCAAGACAGTTTGCACAGAGATTCACCTGTT 1905
Db 1719 CTATTAAACGAAGAATTTAAACACATGAGCTCAGTAAACACACCTGATTCAGGGATCAGT 1778
Qy 1906 ACATTTAAAGAGCCTTTAGAAGCAAAAGAACTAGCTCGGAGCACTTACGAGACCACTG 1965
Db 1779 TTACTTCAATAAAAGTCATCAGAAAGGAAGAAATTTATGCCAAAGACTTTAAAGAACAA 1838
Qy 1966 GATGAAGTGGAGAAAGAACTAGATCAAACTACAGAGATTCATATTTCAATATATCAG 2025
Db 1839 GATGCTCTTTGAAAGAAAGAACTGCATCTAAGCTCTCAGAAATGGAATTCATTTAAACAATCAG 1898
Qy 2026 CTGAAGGAAGTGAAGAAATACACAATAAGCAACAACTCCAGAAAGCAAAAGTCCATGGAG 2085
Db 1899 CTGAAGGAAGTCTAGAAAGAGCTTAAATACACAGCTTAGCCCTTGAACAACTTCATAA 1958
Qy 2086 GCTGAAGCAGCTGAAAACAGAAAGCAAGAAACGAAAGATCATGAATTTAGAAAAACAAA 2144
Db 1959 ATCAACAGTGCACAAATTTGAAGGAAATCGAAAGAAAGAAATTTAGAGCAAAAAA 2017

RESULT 4
US-09-215-681-72
; Sequence 72, Application US/09215681A
; Patent No. 6528253
; GENERAL INFORMATION:
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: Frudakis, Tony N.
; APPLICANT: King, Gordon E.
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSIS
; TITLE OF INVENTION: OF OVARIAN CANCER
; FILE REFERENCE: 210121.463
; CURRENT APPLICATION NUMBER: US/09/215,681A
; CURRENT FILING DATE: 1998-12-17
; NUMBER OF SEQ ID NOS: 310
; SOFTWARE: Fast-Seq for Windows Version 3.0
; SEQ ID NO 72
; LENGTH: 2017
; TYPE: DNA
; ORGANISM: Homo sapien
US-09-215-681-72

Query Match 9.88; Score 507.8; DB 3; Length 2017;
Best Local Similarity 55.88; Pred. No. 7.3e-111;
Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5;

Qy 175 GGGCTCGATTAGCAAGGTAAAGGTAAACAGAACCCATGGCTCAGTTTCAACACCTTTGGT 234
Db 9 GGCTGAGAGCTGCAAGAAAGTCCAGGATCATGTGCTCAGTTTCCACAGCGATGAAT 68
Qy 235 GGCAGCTCGATATCTGGGCCATAAATCTGTAGAGAAAGAGGAGGAGCATGATCAGCAGTTC 294
Db 69 GGAGGGCAAAATATGTGGCTATTACATCTGAAGAACGTACTAAGCATGATAAAGAGTTT 128
Qy 295 CATAGTTTAAAGCAATATCTGGATTTCATTCGTGATCAAGCTAGAAACTTTTTTTTTT 354
Db 129 GATAACCTCAACCTTCAGGAGGTATCATACAGGTGATCAAGCCCGTACTTTTTTCTTA 188
Qy 355 CAATCTGGGTATACCTCAACCTGTTTTTAGCACAGATATGGGCATATAGCTAGCATGAAT 414

Db 189 CAGTCAGGCTGCGGCCCGCGTTTATAGCTGAAATATGGCCCTTATCAGATCTGAACAAG 248
QY 415 GATGGAAGAAATGATCAAGTGGAGTTTCCATAGCTATGAAGCTTATCAAACTGAAGCTA 474
Db 249 GATGGAAGATGGACAGCAAGATTCTCTATAGCTATGAAGCTATCAAGTTAAAGTTG 308
QY 475 CAAGGATATCAGCTACCTCTGTCACCTTCCCTCTGTCATGAAGACAGCAAC--AGTTGCT 531
Db 309 CAGGCGCAACAGCTGCTGATGCTCTCCCTCTCTCTATCATGAACAACCCCTATGTTCTCT 368
QY 532 ATTTCTAGCCACCACATTTGTATGGAGGATCGCCAGCATGCCAGCTTACAGCT 591
Db 369 CCACATAATCTCTGCTCGTTTGGGATGGGAAGCATGCCCAATCTGTCCATTATCAGCCA 428
QY 592 GTTGCTCCAGTGCATGGGATC-----CAAT 618
Db 429 TTGCTCTCAGTTGCACCTATAGCAACACCTTGTCTTCTGCTACTTCAGGGACCAGTAT 488
QY 619 CCAGTTGTTGGAATGCTCCAAACCTTAGTATCTTCTGTTCCACAGCAGCTGTGCCCCC 678
Db 489 CTTCCCTTAATGATGCTGCTCCCTAGTGCCTTCTGTTAGTACATCTCTCAATACCAAT 548
QY 679 CTGGCTAAAGGGGCTCCCTCTTATACAACTCTGCCCTGCAATGCTCTCATCTCGAGCC 738
Db 549 GGAAGCTGCCAGTCTCATCAGCTTTTCAATTCCTTATCTTCTTCAACATTTGCTCAT 608
QY 739 ACATTGCCAAAGATTCTTCTTTAGTAGATCTGGTCCAGGGTCACAACTAAACACATAA 798
Db 609 GCATCATCTTACAGCTGATGGAGGATTTGGTGGTGTAGTAGTACAGAGGCCAG 668
QY 799 TTACAAAGGCGACAGTCATTTGATGTGGCGAGTGTCCACAG-----841
Db 669 TCTCTGATTGATTAGGATCTAGTAGCTCAATCTCCTCAACTGCTTCCCTCTCAGGGAAC 728
QY 842 -----TGGCAGGTGGCTGTCTCAGTCATCAGACTGGAATACAGG 885
Db 729 TCACCTTAAGACAGGGACCTCAGAGTGGCGAGTTCTCCAGCCTTCAAGATTAAAGTATCGG 788
QY 886 CAATTATTCATAGTCATGACAAAATATGAGTGGACACTTAACAGGTGCCCAAGCAAGA 945
Db 789 CAAAATTTAATAGTCTAGCAAAAGCATGAGCGGATACCTCTCAGTTTCAAGCTAGA 848
QY 946 ACTATTTATCAGTCAAGTTTACACAGGCTCAGCTGGCTTCAATATGGAATTTTCT 1005
Db 849 AATGCCCTTCTTCAGTCAAACTCTCTCAAACTCAGTAGCTACTATTGGACTCTGGCT 908
QY 1006 GACATTGATCAAGTGGAAACTTACAGCAGAGGATTTATCTGGCAATGCACCTCAT 1065
Db 909 GACATCGATGGTGACGGACAGTTGAAAGCTGAAGAATTTATTTCTGGCGATGCACCTCACT 968
QY 1066 GATGTAGCTATGCTGGCCAAACCACTGCCACCTGTCTGCTCCAGAAATACATTCACCT 1125
Db 969 GACATGCCCAAGCTTGGACAGCACTACCACTGACGTGTGCTCCGAGCTTGTCCCTCCA 1028
QY 1126 TCTTTTGAAGAGTTTCGATCTGGCAGTGGTATATCTGTATAGCTCAACATCTGTPAGAT 1185
Db 1029 TCTTTTCAAGGGGGAAGCAAGTTGAT-----TCTGTAAATGGAACCTCTGCTTTCATAT 1082
QY 1186 CAGAGCTACCAGAGAACCGATTTTAGAGATGACACACAACTATAGAAAGAAATTA 1245
Db 1083 CAGAAACACAAAGAAGAGCGCT-----CAGAAGAAACTG 1118
QY 1246 CTGTAACTGTTGAAGATAAGAGCGGGAGAACTTTGAACGTGGCAACCTGGAACTGGAG 1305
Db 1119 CCAGTTACTTTTGGAGACAAACGGAAGCCNACTATGACAGAGAAACATGGAGCTGGAG 1178
QY 1306 AAACGAAGCAAGCTCTCTCTGGAACAGCAGCGCAAGGAGCAGAGCGCTTGGCCAGCTG 1365
Db 1179 AAGCGACGCAAGTTGTTGATGGAGCAGCAGAGGGAGGCTGAACGCAAGGCCAGAAA 1238
QY 1366 GAGCGCGGCGACGAGGAGGAGGAGCTGTAGCGCCAGGAGCAAGAGCGCAAAACAAA 1425

Db 1239 GAGAAGGAAGTGGGAGCGGAAACAGAGAGAACTGCAAGACAGAATGGAGAAGCAG 1298
QY 1426 CTGGAACCTGGAGAAGCAACTGGAAGAGCAGCGGGAGCTAGAACGCGCAGAGAGAGGAG 1485
Db 1299 CTGAGTTGGAGAAACGCTTGGAGAAACAGAGAGAGCTGGAGAGACAGCGGAGGAAGAG 1358
QY 1486 AGGAGAAAGAAATTTAGAGAGCGGAGAGCTGCNAAACGGGAACCTTGAAGGCAACGACAA 1545
Db 1359 AGGAAAGAGGATAGAAAGACGAGAGCGAAGCAAAACAGGAGCTTTGAGAGACAAACGCGCT 1418
QY 1546 CTGTAGTGGGAACGGAATCGAAGGCAAGAACTTACTATAATCAAAAGAAACAAAGAAACAAGAG 1605
Db 1419 TTAGAAATGGGAAGAGCTCGTCCGACAGAGCTGCTCAGTCAGAAAGCAGGGAACAAGAA 1478
QY 1606 GACATAGTTGTACTGAAAAGCAAGAAAGAAAGCTTTTGGAAATTTGAATTTAGAAGCTCTAAAT 1665
Db 1479 GACATTTGTGAGCTGAGCTCCAGAAAGAAAGTCTCCACCTGGAACTGGAGCAGTGAAT 1538
QY 1666 GATAAAAGCATCAACTAGAAGGGAACCTTCAAGATATCAGATGTCGATTTGACCAACCAA 1725
Db 1539 GGAAGAAATCATCAGCAGATCTCAGGCGAGACTACAAGATGTCCAAATCAGAAAGCAAAACAAA 1598
QY 1726 AGGCAAGAAATTTAGAGCAGCAAAACAAATCTTAGAGAGTTGAGAATTTGCCGAAATCAACCAT 1785
Db 1599 AGACTGAGCTAGAAGTTTGGATAAACAGTGTGACCTGGAAATTTATGMAATCAACAA 1658
QY 1786 CTACAGCAACAAATTAAGGAATCTCAGCAAAATGTTTGGAAAGACTTTATTTCCAGAAAAACAG 1845
Db 1659 CTTCAACAAGAGCTTTAAGCAATATCAAAATAAGCTTTATCTATCTGCTCCTCGAGAAGCAG 1718
QY 1846 ATACTCAATGACCAATTAACAACTTTCAGCAGACACAGTTTGCACAGAGATTCACCTGTT 1905
Db 1719 CTATTAAACGAAAGAAATTTAAACCAATGCAGCTCAGTAAACACCTGATTCAGGGATCAGT 1778
QY 1906 ACACTTAAAGAGCTTTAGAAGCAAAAGAACTAGCTCGGCGACACTACGAGACCAACTG 1965
Db 1779 TTACTTCATAAAAGTCTATCAGAAAGAGAGAAATTTGCCAAAGACTTTAAGAACAAATTA 1838
QY 1966 GATGAAGTGGAGAAAGAACTAGATCAAACTTACAGGAGATTTGATTTTCAATTAATCAG 2025
Db 1839 GATGCTCTTGAAGAAAGAACTGCATCTAAGCTCTCAGAAATGGATTTCATTTAAACAATCAG 1898
QY 2026 CTGAAGGAACTAAGAAATACACAACTTAAGCAACAACTCCAGAGCAAAAGTCCATGGAG 2085
Db 1899 CTGAAGGAACTCAGAGAAAGCTTATATACACAGCAGTTAGCCCTTTGAACAACTTCATAAA 1958
QY 2086 GCTGAACCACTGAAACAGAAAGAAACAGAAACAAAGATCATAGAAATTTAGAAAAACAAA 2144
Db 1959 ATCAACCTGACAAATTTGAAGAAATCGAAGAAAGAAATTTAGACCAAAAGAAAAAAA 2017

RESULT 5

US-09-216-003A-72

; Sequence 72, Application US/09216003A

; Patent No. 6670463

; GENERAL INFORMATION:

; APPLICANT: Mitcham, Jennifer L.

; APPLICANT: Frudakis, Tony N.

; APPLICANT: King, Gordon E.

; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY OF OVARIAN CANCER

; FILE REFERENCE: 210121.462

; CURRENT APPLICATION NUMBER: US/09/216,003A

; CURRENT FILING DATE: 1998-12-17

; NUMBER OF SEQ ID NOS: 310

; SOFTWARE: PatentIn Ver. 2.0

; SEQ ID NO 72

; LENGTH: 2017

; TYPE: DNA

; ORGANISM: Homo sapiens

US-09-216-003A-72

Query Match 9.8%; Score 507.8; DB 3; Length 2017;

Best Local Similarity 55.8%; Pred. No. 7.3e-111;

Matches	1138;	Conservative	0;	Mismatches	802;	Indels	99;	Gaps	57;
175	QY	GGCGTCGATTAGCAAGTAAAAAGTAAACAGAACCATGGCTCAGTTTCCAAACACCTTTTGGT	234						
9	Db	GGCTGAGAGCTGCAAGAAAGTCCAGGATCATGATGGCTCAGTTTCCCAACAGCGATGAAT	68						
235	QY	GGCAGCTGGATATCTGGGCCATTAACCTGTAGAGGAAGAGCGAAGCATGATCAGCAGTTTC	294						
69	Db	GGAGGGCCAAATATGTGGGCTATTACATCTGAAGAACGCTACTAAGCATGATATAACAGTTT	128						
295	QY	CATAGTTTAAAGCCAATATCTGGATTCAATTACTGTGGTGATCAAGCTAGAAACTTTTTTTT	354						
129	Db	GATAACCTCAAACTTCAGGAGGTACATAACAGGTGATCAAGCCGTACTTTTTTCCCTA	188						
355	QY	CAATCTGGGTTACCTCAACCTGTTTTAGCACAGATATGGCACTAGCTGACATGAATAAT	414						
189	Db	CAGTCAGGTCGCGGCCCGGTTTTAGCTGAATAATGGCCCTTATCAGATCTGAAACAAG	248						
415	QY	GATGGAAGTAATGATCAAGTGAGTTTTCATAGCTATGAAGCTTATCAACTGAAGCTA	474						
249	Db	GATGGGAAGATGGACAGCAAGAGTTTCTCTATAGCTATGAAACTCATCAAGTTTAAAGTTG	308						
475	QY	CAAGGATATCAGTACCTCTGCACTTCCCTCTCATGAAACAGCAAC--AGTTGCT	531						
309	Db	CAGGGCCAAACAGCTGCTGTATGTCCTCCCTCATGAAACCAACCCCTATGTTCTCT	368						
532	QY	ATTCTTAGCGCACAGCATTTGGTATGGGAGGTATGCCAGCATGCCACGGCTTACAGCT	591						
369	Db	CCACTAACTCTGCTCGTTTTGGGATGGGAAGCATGCCCAATCTGTCCATTATCAGCCA	428						
592	QY	GTTGCTCCAGTGCCAAATGGGATC-----CAATT	618						
429	Db	TTGCCCTCCAGTTCGACCTATAGCAACACCCCTTGTCTTCTGCTACTTTCAGGGACCAGTA	488						
619	QY	CCAGTTGTTGGATGTCCTCAACCTTAGTATCTTCTGTTCCCAACAGCAGCTGCCCCC	678						
489	Db	CCTCCCCTAATGATGCTGCTCCCCAGTGCCTTCTGTTAGTACATCTCTATTACCAAT	548						
679	QY	CTGGCTAAAGGGGCTCCCCCTGTTTATACAACTCTGCTGCTATGCTATCTCTCAGCC	738						
549	Db	GGNACTGCCAGTCTCATTCAGCCTTTATCCATTCCCTTATCTTCTTCAACATTGCTCAT	608						
739	QY	ACATTGCCAAAGAGTTCTTCTCTTTAGTAGATCTGGTCCAGGGTCACAACCTAAACCTAA	798						
609	Db	GCATCATCTTTACAGCCTGATGATGGGAGGATTTGGTGTCTAGTATCCAGAAGGCCAG	668						
799	QY	TTACAAAGGCACAGTCATTTGATGTGGCAGTGTCCACCG-----841							
669	Db	TCTCTGATTGATTTAGAGCTTAGTAGCTCAACTTCTCTCAACTGCTTCCCTCTCAGGGAAC	728						
842	QY	-----TGGCAGAGTGGGCTGTTTCTCAGTCATCAAGACTGAAATACAGG	885						
729	Db	TCACCTAAGACAGGGACCTCAGAGTGGGAGTTTCTCAGCCITCAAGATTAAAGTATCGG	788						
886	QY	CAAATTATCAATAGTCATGACAAACATPATCAGTGGACACTTAAACAGTCCCCAAGCAAGA	945						
789	Db	CAAAATTTAATAGCTTAGACAAAGGCATCAGCGGATACCTCTCAGGTTTTCAAGCTAGA	848						
946	QY	ACTATTCTTATGCAGTCAAGTTTACACAGGCTCAGCTGGCTTCAATATGGATCTTTCT	1005						
849	Db	AATGCCCTTCTCAGTCAAAATCTCTCTCAAACTCAGCTAGCTACTATTTTGGACTCTGGCT	908						
1006	QY	GACATTGATCAAGATGAAAACCTTACAGCAGAGGAATTTATCTGGCAATGCACTCATTT	1065						
909	Db	GATCATGNTGGTGACGGACAGTTGAAAGCTGAAGAAATTTATCTGGCGATGCACCTCACT	968						
1066	QY	GATGTAGTATGTCTGGCCCAACCACTGCCACTGTCTCTGCTCCAGAAATACATTCACCT	1125						
969	Db	GACATGSCCAAGCTGGACAGCCACTACCACTTGACGTTGCTCCCGAGCTTGTCCCTCCA	1028						
1126	QY	TCCTTTTGAAGAGTTTCGATCTGGCAGTGGTATATCTGTCTATAGCTCAACATCTGTAGAT	1185						
1029	Db	TCCTTTACAGGGGGAACGAAGTTGAT-----TCTGTTAATGGAATCTGCTCTCATAT	1082						

1186	Qy	CAGAGCGCTACCAGAGAAACCGAGTTTGTAGAGATGACACAACTTGTAGAAAAGAAATTA	1244
1187			
1083	Db	CAGAAAACAACAAGAGAGAGCGCT-----CAGAAGAAACTG	1118
1246	Qy	CTGTAAACGTTTGAACATAAGAAAGCGGGAGAACTTTGAAACGTGGCAACCTGGAACTGGAG	1305
1119	Db		
1119	Db	CCAGTTTACTTTTGAGAGCAAAACGGAAGCCNACTATGAACGAGGNAACATGGAGCTGGAG	1178
1306	Qy	AAACGAAGCGAAGCTCTCTCTGGAAACAGCAGCGCAAGGACGAGAGCGCCTGGGCCACGCTG	1365
1179	Db	AAGCGACGCCAAGTGTGTGATGGAGCAGCAGCAGAGGGAGGCTGAAGCGCAAGGCCACGAAA	1238
1366	Qy	GACCGGGCGGACAGAGAGGAGGAGCGCTGTAGCGCCAGGACGCAAGAGCGCAAAAGACAA	1425
1239	Db	GAGAAAGGAGAGTGGGAGCGGAAACAGAGAGAACTGCAAGGACAGAACTGGNAGAGCAG	1298
1426	Qy	CTCGAACTGGAGAAAGCAACTTGGAAAAACGCGGGAGCTAGAAACGGCAGAGAGAGGAGGAG	1485
1299	Db	CTGGAGTTGGAGAAAACGCTTGGAGAAACAGAGAGAGCTGAGAGACAGCGGGAGGAAGAG	1358
1486	Qy	AGGAGGAAAGAAATTTGAGAGCGCAGAGGCTTGCAAAAACGGGAACTTTGAAAGGCAACGACAA	1545
1359	Db	AGGAGAAAGAGATAGAAACAGCAGAGGCGACAAAACAGGAGCTTGAGAGACAAACGCGCT	1418
1546	Qy	CTTGAGTGGGAAACGGAACTCGAAGGCAAGAACTACTAAATCAAGAAACAAAGAACCAAGAG	1605
1419	Db	TTAGAAATGGGAAAGACTCCGTCGGCAGGAGCTGCTCAGTCAGAGAACACGGGAAACAAGAA	1478
1606	Qy	GACATAGTTGTACTGAAAGCAAAAGAAAAGACTTTTGGAAATTTGAAATTTAGAAGCTCTAAAT	1665
1479	Db	GACATTTGTCAGGCTGAGCTCCAGAAAGNAAGTCTCCACCTGGAACTGGAAGCAGTGAAT	1538
1666	Qy	GATAAAAGCATCAACTAGAAAGGGAACCTTCAAGATATCAGATGTGCGATTGACCAACCCAA	1725
1539	Db	GGAAACATCAGCAGATCTCAGGCAGACTACAAGATGTCCAAATCAGAAAGCAAAACAA	1598
1726	Qy	AGGCAAGAAATTTGAGAGCAAAACAAATCTAGAGAGTTGAGATTCGCGAAATCAACCCAT	1785
1599	Db	AAGACTGAGCTAGAAAGTTTTTGGATATAACAGTGTGACCTGGAAATATATGSAATCAAAACAA	1658
1786	Qy	CTACAGCAACAAATTTACAGGAATCTCAGCAAAATGCTTGGAAAGACTTATTCAGAAAAACAG	1845
1659	Db	CTTCAACAGAGCTTAAGGAATATCAAAATAGCTTATCTATCTGTGCTCTGTGAGAGCAG	1718
1846	Qy	ATACTCAATGACCAATTTAAACAAAGTTTCAGCAGAACAGTTTGGCACAGAGATTCATCTGTT	1905
1719	Db	CTATTAAACGAAGAAATTAATAACATGCACCTCAGTAACACACCTGATTCAGGGATCAGT	1778
1906	Qy	ACACTTAAAGAGCCTTAGAAGCAAAAGAACTAGCTCGGACGACNCTCAGAGACCAACTG	1965
1779	Db	TTACTTCATAAAAGTCAATCAGAAAAGGAGAAATTAATGCCAAAAGACTTAAAGAAACAAATTA	1838
1966	Qy	GATGAAGTGGAGAAAGAAATCTAGATCAAAACTACAGAGAGTTGATATTTTCAATAATACAG	2025
1839	Db	GATGCTCTTGAAAAAGAAATCTGCATCTTAAGCTCTCAGAAATGGATTCATTTTACAAATCAG	1898
2026	Qy	CTGAAGGAACTTAAGAGAAATACACAAATAAGCAACAACTCCAGAGCAAAAAGTCCATGGAG	2085
1899	Db	CTGAAGGAACTCAGAGAAAGCTATAATACAGCAGTGTAGCCCTTGAACAACTTCATAAA	1958
2086	Qy	GCTGAACGACTGAAACAGAAAGAAACAAGACGAAAGATTCATAGAAATTTAGAAAAACAAA	2144
1959	Db	ATCAAACTGTGCAAAATTTGAAGGAAATCGAAAGAAAAAGATTTAGAGCAAAAAAANAANA	2017

RESULT 6
US-09-667-857-72
; Sequence 72, Application US/09667857
; Patent No. 6699664
; GENERAL INFORMATION:
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: King, Gordon E.

APPLICANT:	Algate, Paul A.	APPLICANT:	Fling, Steven P.	APPLICANT:	Reiter, Marc W.	APPLICANT:	Panger, Gary Richard	APPLICANT:	Reed, Steven G.	APPLICANT:	Vedvick, Thomas S.	APPLICANT:	Carter, Darrick	TITLE OF INVENTION:	COMPOSITIONS AND METHODS FOR THE THERAPY AND	TITLE OF INVENTION:	DIAGNOSIS OF OVARIAN CANCER	FILE REFERENCE:	210121.462C5	CURRENT APPLICATION NUMBER:	US/09/667,857	CURRENT FILING DATE:	2000-09-20	NUMBER OF SEQ ID NOS:	455	SOFTWARE:	FastSeq for Windows Version 3.0	SEQ ID NO	72	LENGTH:	2017	TYPE:	DNA	ORGANISM:	Homo sapien	US-09-667-857-72
Query Match	9.8%;	Score	507.8;	DB	3;	Length	2017;	Best Local Similarity	55.8%;	Pred.	No. 7.3e-111;	Matches	1139;	Conservative	0;	Mismatches	802;	Indels	99;	Gaps	5;															
175	GGCGTCGATTAGCAAGGTAAAGTAAACAGAAACATGGCTCAGTTTCCAAACCTTTTGGT	234	9	GGCTGAGAGCTGCAAGAAAGTACAGATCATGATGGCTCAGTTTCCACAGCGATGAAT	68																															
235	GGCAGCTGGATCTGGGCCATACTGTAGAGGAAGAGCGAAGCATGATCAGCGTTTC	294	69	GSAGGGCCAAATATGGGGCTATTACATCTGAAGAACGCTACTAAGCATGATATAACAGTTT	128																															
295	CATAGTTTAAAGCCAAATATCTGGATTCATTACTTGGTGATCAAGCTPAGAAACTTTTTTTTT	354	129	GATPACTCAACCTTCAGAGGTTTACATAACAGGTGATCAAGCCGCTACTTTTTTCTTA	188																															
355	CAATCTGGGTTACCTCAAACCTGTTTTCAGACAGATATGGGCACTAGCTGACATGAATAT	414	189	CAGTCAGGCTGCGGCCCGGTTTTAGCTGAAATATGGGCTTATCAGATCTGAAACAAG	248																															
415	GATGGGAAGATGATCAAGTGGAGTTTCCATAGCTATGAACCTTATCAAACTGANGCTA	474	249	GATGGGAAGATGACAGCAAGAGTTTCTATAGCTATGAAACTCATCAAGTTAAAGTTG	308																															
475	CAAGGATATCAGCTACCTCTGCACCTTCCCTCTCATGAAACAGCAACC--AGTTGCT	531	309	CAGGCCAACAGCTGCTGTATGCTTCCCTCTCATGAAACAGCAACCCTTATGTTCTCT	368																															
532	ATTCTAGCGCACAGCATTTGGTATGGAGGTATGCGCAGCATGCCACCGCTTACAGCT	591	369	CCACTAATCTCTGCTCGTTTGGGATGGGAAGCATGCCCAATCTGTCCATTATCAGCCA	428																															
592	GTGCTCCAGTGCATGGGATC-----CAATT	618	429	TTGCGCTCCAGTTGCACCTATAGCAACCCCTTGTCTTCTGTCTTTCAGGACCAAGTAT	488																															
619	CCAGTTGTTGGAATGTCCTCAACCTTAGTATCTTCTGTTCCACAGCAGCTGTGCCCC	678	489	CTTCCCTTAATGATGCTGCTCCCTTAGTGCCCTTCTGTTAGTACATCTCATTTACCAAT	548																															
679	CTGGCTAACGGGGCTCCCTGTTATACAACTCTGCTGCAATTTGCTCATCTGCGAGCC	738	549	GGAACCTGCAGTCTCATTCAGCCCTTTATCCATTTCTTATTCTTCTTCAACATTCGCTCAT	608																															
739	ACATTGCCAAAGAGTTCTCTTTTAGTATGATCTGGTTCAGGGTCAACAATAACACTAAA	798	609	GCATCATCTTACAGCTGATGATGGGAGGATTTGGTGTGTAGTATCCAGAAAGGCCAG	668																															
799	TTACAAAGGCCACAGTCATTTGATGTCGGCAGTGTCCTCCACAG-----	841	669	TCTCTGATTGATTAGGATCTAGTAGCTCAACTTCTCTCAACTGCTTCCCTCTCAGGGAAC	728																															
842	-----TGGCAGAGTGGGCTGTTCTCTCAGTCATCAAGACTGAAATACAGG	885																																		

Db 1779 TTACTTCATAAAAGTCATCAGAAAAGGAGAAATATATGCCAAGACCTTAAAGAACAAATTA 1838
Qy 1966 GATGAAGTGGAGAAAGAACTAGATCAAACTACAGGAGATGATATTTTCAATATATCAG 2025
Db 1839 GATGCTCTTGAAGAAAGAACTGATCTAAGCTCTCAGAAATGATTTCAITTAACAATCAG 1898
Qy 2026 CTGAAGGAACCTAGAGAAATACAAATAGCAACAACTCCAGAAAGCAAAAGTCCATGGAG 2085
Db 1899 CTGAAGGAACCTAGAGAAAGCTATATACACAGAGTTAGCCCTTGAACAACTTCATATAA 1958
Qy 2086 GCTGAACGACTGAAACAGAAAGAAACAAAGAACGAGAAAGATCATAGAAATAGAGAAACAAA 2144
Db 1959 ATCAAAGCTGACAAATTTGAAGAAATCGAAGAAAGAAAGATTAGAGCAAAAGAAAAA 2017

RESULT 7

US-10-198-053-72

; Sequence 72, Application US/10198053

; Patent No. 6858710

; GENERAL INFORMATION:

; APPLICANT: Bangur, Chaitanya S.

; APPLICANT: Retter, Marc W.

; APPLICANT: Fanger, Gary R.

; APPLICANT: Hill, Paul

; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

; FILE REFERENCE: 210121.462C9

; CURRENT APPLICATION NUMBER: US/10/198,053

; CURRENT FILING DATE: 2002-07-17

; NUMBER OF SEQ ID NOS: 624

; SOFTWARE: FastSeq for Windows Version 4.0

; SEQ ID NO 72

; LENGTH: 2017

; TYPE: DNA

; ORGANISM: Homo sapiens

US-10-198-053-72

Query Match 9.8%; Score 507.8; DB 3; Length 2017;

Best Local Similarity 55.8%; Pred. No. 7.3e-111;

Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5;

Qy 175 GCGCTCGATTAGCAAGCTAAAGTAAACAGAACCATGGCTCAGTTTCCACACACCTTTTGGT 234
Db 9 GCGTGAGAGCTGCAAGAAAGAGTCAGGATCATGATGGCTCAGTTTCCACAGCGATGAAT 68
Qy 235 GCGAGCCTGGATATCTGGGCAATACTGTAGAGGAAAGAGCGAAGCATGATCAGCAGTTC 294
Db 69 GGAGGGCCAATATGTGGCTATTACATCTGAAGAACGTACTAAGCATGATAAACAGTTT 128
Qy 295 CATAGTTTAAAGCAATATCTGGATTCATTCTAGTGTATCAGTGAAGAACTTTTCTTTT 354
Db 129 GATAACCTCAAACTTCAGGAGGTTACATAACAGGTGATCAAGCCCGTACTTTTCTCTA 188
Qy 355 CATCTGGGTACCTCAACCTGTTTATAGCAGATATGGGCATCTAGCATGAATAT 414
Db 189 CAGTCAGTCTGCGGCGCGGTTTATAGCTGAATATGGGCTTTATCAGATCTGAACAG 248
Qy 415 GATGGAAGAAATGGAATCAAGTGGAGTTTTCATAGCTATGAACCTTATCAAACTCAAGCTA 474
Db 249 GATGGGAAGATGGACGCAAGAGTCTCTATAGCTATGAACCTCAATCAAGTTAAAGTTG 308
Qy 475 CAAGGATATCAGCTACCTCTGCACTTCCCTCTGATGAAGAAAGCAAGCAAC---AGTTGCT 531
Db 309 CAGGCGCAACAGCTGCTGTAGTCTCCCTCTATCATGAAGCAACCCCTTATGTTCTCT 368
Qy 532 ATTTCTAGCGCACCGAGATTTGGTATGGAGGTATCGCCAGCATGCCACCGCTTACAGCT 591
Db 369 CCACATAATCTCTGCTCGTTTGGGATGGAAAGCATGCCAATCTGTCCATTCATCAGCCA 428
Qy 592 GTTGCTCAGTGCCAAATGGGATC-----CATT 618
Db 429 TTGCTCCAGTTGCACCTATAGCAACACCCCTTGTCTTCTGCTACTTTCAGGAGCAGTATT 488

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Db      1539  GGAAGAAATCATCAGCAGATCTCAGCAGACTACAAGATGTCCAATCAGAAAGCAACACAA 1598
Qy      1726  AGCGAAGAAATTTGAGAGCACAACAAATCTAGAGAGTTGAGAAATTCGCCAAATCACCCAT 1785
Db      1599  AAGACTGAGCTAGAAGTTTGGATAAACAGTGTGACCTGGAAATTTATGGAATTCAAACAA 1658
Qy      1786  CTACAGCAACAAATTCAGAGAAATCTCAGCAAAATGCTTTGGAAGACTTTATTCAGAAAAACAG 1845
Db      1659  CTTCAACAAGAGCTTAAGGAATATCAAAATAGCTTTATCTATCTGGTCCCTGAGAAGCAG 1718
Qy      1846  ATACTCAATGACCAATTAACAAAGTTTCAGCAGAACAGTTTGACACAGAAATTCACATTGTT 1905
Db      1719  CTATTAAACGAAAGAAATTAACAAACATGCGAGCTCAGTAAACACACCTGATTCAGGGATCAGT 1778
Qy      1906  ACACCTTAAAGAGCCCTTGAAGCAAAAGAACTAGCTCGGCAGCACCTCAGAGACCAACTG 1965
Db      1779  TTACTTCAATAAAGTCATCAGAAAGGAAGAAATTTATGCGCAAGACTTAAAGAACAAATTA 1838
Qy      1966  GATGAAGTGGAGAAAGAACTAGATCAAAACTACAGGAGATTCATATTTTCAATAATCAG 2025
Db      1839  GATGCTCTTTGAAAAAGAAACTGCATCTAAGCTCTCAGAAATGGAATTCATTTAAACAATCAG 1898
Qy      2026  CTGAAGGAATCAGAGAAATACAAATAGCAACAACTCCAGAAAGCAAAAGTCCATGGAG 2085
Db      1899  CTGAAGGAATCAGAGAAAGCTATATACACAGCAGTTAGCCCTTTGAAACAACCTTCATATAA 1958
Qy      2086  GCTGAACGACTCAAAACAGAAAGAAACAAAGAAAGCAAGAACTATAGAAATTAGAAAAACAAA 2144
Db      1959  ATCAAAAGTCACAAATGAAGGAAATCGAAAGAAAGAAATTAGAGCAAAAGAAAAA 2017
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RESULT 8

US-09-827-271-72

; Sequence 72, Application US/09827271

; Patent No. 6962980

; GENERAL INFORMATION:

; APPLICANT: Retter, Marc W.

; APPLICANT: Fanger, Gary R.

; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

; TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER

; FILE REFERENCE: 210121.462C6

; CURRENT APPLICATION NUMBER: US/09/827,271

; CURRENT FILING DATE: 2001-04-04

; NUMBER OF SEQ ID NOS: 461

; SOFTWARE: FastSeq for Windows Version 3.0

; SEQ ID NO 72

; LENGTH: 2017

; TYPE: DNA

; ORGANISM: Homo sapien

US-09-827-271-72

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Query Match          9.8%; Score 507.8; DB 3; Length 2017;
Best Local Similarity 55.8%; Pred. No. 7.3e-111;
Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5;

Qy      175  GCGTCGATTAGCAAGTAAAAGTAAACAGAACCATGGCTCAGTTTCCAAACACCTTTTGGT 234
Db      9    GCGTGAGAGCTGCAAGAAAGTCAGGATCATGTGGCTCAGTTTCCCACAGCGATGAAT 68
Qy      235  GCGAGCCTGGATATCTGGGCCATAACTGTAGAGGAAGAGCGNAGCATGATCAGCGATTC 294
Db      69  GGAGGGCCAAATATGTGGGCTATTACATCTGAAGAACGTAATAAGCATGATAACAGTTT 128
Qy      295  CATAGTTTAAAGCCAAATATCTGGATTCATTACTGGTGATCAAGCTAGAAACTTTTTTTTT 354
Db      129  GATAACCTCAAACTTCAGAGGTTACATAACAGGTGATCAAGCCCGTACTTTTTTCCTA 188
Qy      355  CAATCTGGGTTACCTCAACCTGTTTTTACACAGATATGGGCACTAGCTGACATGAATAAT 414
Db      189  CAGTCAGGTCTGCGGCCCGGTTTTTAGCTGAAATATGGGCTTTATCAGATCTGAACAAG 248
Qy      415  GATGGAAGATGGATCAAGTGGAGTTTTTCCATAGCTATGAAACTTATCAAACTGAAGCTA 474
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Db      249  GATGGGAAGATGGACAGCAAGAGTTCTATAGCTATGAAACTCATCAAGTTAAAGTTG 308
Qy      475  CAAGGATATCAGCTACCCCTCTGCACCTCCCTCTGTCATCGTCAAGAAACAGCAACC---AGTTGCT 531
Db      309  CAGGCCCAACAGCTGCTGTAGTCTCTCTCTATCATGAACAACCCCTATGTTCTCT 368
Qy      532  ATTTCTAGCGCACCAAGCATTTTGGTATGGAGGATATCGCAGCATGCCACGGCTTACAGCT 591
Db      369  CCACATAATCTCTGCTCGTTTTGGGATGGGAAGCATGCCCAATCTGTCCATTCATCAGCCA 428
Qy      592  GTTGCTCCAGTGCCTAATGGGATC-----CATT 618
Db      429  TTGCTCCAGTTGCACCTATAGCAACACCTTGTCTTCTGTACTTTCAGGGAGCCAGTATT 488
Qy      619  CCAGTTGTTGGAATCTCTCCAAACCTAGTATCTCTCTGTTCCACAGCAGCTGTGCCCCCC 678
Db      489  CCTCCCTTAATGATGCTGCTCCCTAGTGCCTTCTGTTAGTACATCCTCATTACCAAT 548
Qy      679  CTGGCTAACGGGGCTCCCTCTGTTATACAACTCTGCTGCAATTTGCTCATCCTTGAGCC 738
Db      549  GGAACCTGCAGTCTCATTCAGCCTTTATCCATTTCTTCTTCAACATTGGCTCAT 608
Qy      739  ACATTTGCCCAAGAGTTCTTCTTTAGTAGATCTGGTCCAGGGTCAACAATAACAATAA 798
Db      609  GCATCATCTTACAGCTCATGATGGGAGGATTTGGTGGTGTAGTATCCAGAAAGGCCAG 668
Qy      799  TTACAAAAGGCACAGCTCATTTGATGTGCCAGTGTCCACCCAG----- 841
Db      669  TCTCTGATTGATTAGGATCTAGTAGCTCAACTTCTCAACTGCTTCCCTCTCAGGGAAC 728
Qy      842  -----TGGCAGAGTGGGCTGTTCTCAGTCATCAAGACTGAAATACAGG 885
Db      729  TCACCTAAGACAGGGACCTCAGAGTGGGCGAGTTCTCAGCCTTCAAGATTAAAGTATCGG 788
Qy      886  CAATTTATTCAATAGTCATGACAAAATATAGTGGACACTTAAACAGGTCCCAAGCAAGA 945
Db      789  CAAAAATTTAATAGTCTAGACAAAGCATGAGCGGATACCTCTCAGGTTTTTCAAGTAGA 848
Qy      946  ACTATTTCTATGCACTCAAGTTTACACAGGCTCAGCTGGCTTCAATATGGAATCTTTCT 1005
Db      849  AATGCCCTTCTTCACTCAAACTCTCTCAAACTCAGCTAGCTACTATTTGGACTCTGGCT 908
Qy      1006  GACATTTGATCAAGATGGAAAACTTACAGAGAGGAATTTATCCTGGCAATGCACTCAT 1065
Db      909  GACATCGATGGTGAACGAGCTGAAAGCTGAAGAAATTTATTTCTGGCGATGCACTCACT 968
Qy      1066  GATGTAGCTATGTCTGGCCCAACCACTGCCACCTGCTCTGCTCCAGAAATACATTCACCT 1125
Db      969  GACATGGCCAAAGCTGGACAGCCCTACCACTGACGTGGCTCCCGAGCTGTGCTCTCA 1028
Qy      1126  TCTTTTAGAAGAGTTTCGATCTGGCAGTGGTATATCTGTATAGCTCAACATCTGTAGAT 1185
Db      1029  TCTTTTACAGGGGGAAAGCAAGTTGAT-----TCTGTTAATGGAACCTGCTTTCATAT 1082
Qy      1186  CAGAGGCTACAGAGGAACCGATTTTAGAAGATGAACAACAATTAGAAAAAGAAATTA 1245
Db      1083  CAGAAAAACACAGAAAGAGAGCCT-----CAGAAAGAAACATG 1118
Qy      1246  CCTGTTAAACGTTTGAAGATAAGAAAGCGGAGAACTTTTGAACGTGGCAACCTCGAACTGGAG 1305
Db      1119  CAGTTTACTTTTGGAGACAAACGGAAGCAACTATGAAACAGGGAACATGAGAGCTGGAG 1178
Qy      1306  AAAAGGAAGGCAAGCTCTCTCTGGAAACAGCAGCGCAAGAGGAGAGAGCGCTGCGCCAGCTG 1365
Db      1179  AAGCGACGCCAAGTGTGTATGGAGCAGCAGCAGAGAGGAGGCTGAAACGCAAGCCAGAAA 1238
Qy      1366  GAGCGGGCGGAGCAGAGGAGGAGCGGTGAGCGCCAGGAGCAGAGCGCAAGGCAAGACAA 1425
Db      1239  GAGAAAGGAAGTGGGAGCGGGAACAGAGAGAACTGCAAGAGACAAAGATGGAAGAGCAG 1298
Qy      1426  CTGGAACCTGGAGAAGCAACTGGAAGAGCAGGGGAGCTAGAACGCGCAGAGAGAGGAGGAG 1485
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Db 1299 CTGGAGTTGGAGAAACGCTTTGGAGAAACAGAGAGAGCTGGAGAGACAGCGGGAGGAGAG 1358
Qy 1486 AGGAGGAGAAATTCAGAGGCGAGAGGCTTGCAAAAACGGGAACTTGAAAGGCAACAGCAAA 1545
Db 1359 AGGAGAAAGGAGATAGAAAGACGAGAGGCGAGCAAAAACAGGAGCTTGAGAGACAAACGCCGT 1418
Qy 1546 CTTGAGTGGGAACGGAATCGAAGGCAAGAACTACTAAATCAAGAAACAAAGAAACAGAGAG 1605
Db 1419 TTAGAATGGGAAGACTCCGCTCGGAGGAGCTCTCAGTCAGAGAGACCGAGGAAACAGAA 1478
Qy 1606 GACATAGTTGCTACTGAGAAACAAAGAAAGAAAGACTTTTGGAAATTTGAAATTTAGAAGCTCTAAAT 1665
Db 1479 GACTTGTGAGCTGAGCTCCAGAAAGAAAGCTCTCACTGGAACCTGGAAGCAGTGAAT 1538
Qy 1666 GATAAAAGCATCAACTAGAGGGAATCTTCAAGATATACAGATGTGATTTGACCAACCCAA 1725
Db 1539 GGAAGAAACATCAGCAGATCTCAGGAGACTACAAGATGTCCAAATCAGAAAGCAACACAA 1598
Qy 1726 AGCAAGAAATTCAGAGCACAAACAAATCTAGAGAGTTGAGAGTTGCGGAATTCGGAATCACCCT 1785
Db 1599 AAGACTGAGCTAGAAGTTTGGATAACAGTGTGACCTGGAATTTATGGAATTCACAA 1658
Qy 1786 CTACAGCAAAATTCAGAGAACTCTCAGCAAAATCTTGGAAAGCTTAATTCAGAAAAACAG 1845
Db 1659 CTTCAACAGAGCTTAAGGATATCAAAATAGCTTATCTATCTGGTCCCTGAGAGCAG 1718
Qy 1846 ATACTCAATGACCAATTTAAACAAAGTTTCAGCAAGAACTTTGACAGAGATTCACCTGTT 1905
Db 1719 CTATTAACGAAAGAAATTTAAACATGCAGCTCAGTAAACACACCTGATTCAGGGATCAGT 1778
Qy 1906 ACATTAAGAGCCTTAGAGCAAGAAAGAACTAGCTCGGAGCACTTACGAGACCAACTG 1965
Db 1779 TTACTTCATAAAAGTCAACAGAAAGAAAGAAATTTATGCCAAAGACTTAAAGAACAAATTA 1838
Qy 1966 GATGAAGTGGAGAAAGAACTAGATCAAAACTACAGAGAGTTGATATTTTCAATAATCAG 2025
Db 1839 GATGCTTTGAAAAGAAAGAACTGCATCTAAGCTCTCAGAAATGGATTCATTTAACAATCAG 1998
Qy 2026 CTGAAGAACTAAGAGAAATACAAATAGCAAACTCCAGAGCAAGAAAGTCCATGAGG 2085
Db 1899 CTGAAGAACTCAGAGAAAGCTATAATACAGAGCAGTTAGCCCTTGAACTTCAATAA 1958
Qy 2086 GCTGAAGCTAGAAACAGAAAGAAACAGACAGAAAGATCATAGAAATTTAGAAACAA 2144
Db 1959 ATCAACGTGACAAATTTGAGGAATTCGAAAGAAAGAAAGATTTAGAGCAAAAAA 2017

RESULT 9
US-09-513-999C-27927
; Sequence 27927, Application US/09513999C
; Patent No. 6783961
; GENERAL INFORMATION:
; APPLICANT: Dumas Milne Edwards, J.B.
; APPLICANT: Duclert, A.
; APPLICANT: Giordano, J.Y.
; TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
; PATENT NO. 6783961
; FILE REFERENCE: 59, US2, REG
; CURRENT APPLICATION NUMBER: US/09/513,999C
; CURRENT FILING DATE: 2000-02-24
; PRIOR FILING DATE: 1999-02-26
; NUMBER OF SEQ ID NOS: 36681
; SOFTWARE: Patent.pm
; SEQ ID NO 27927
; LENGTH: 174
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-513-999C-27927

Query Match 3.3%; Score 174; DB 3; Length 174;
Best Local Similarity 100.0%; Pred. No. 1.2e-31;
Matches 174; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 3936 TTCCCCACCTTTTGCACAGGTGCTTTCAATAGTTTTTAAAAATATATTTTAAATATATATTT 3995
Db 1 TTCCCCACCTTTTGCACAGGTGCTTTCAATAGTTTTTAAAAATATATTTTAAATATATATTT 60
Qy 3996 TAGCTTTTAAATAAACAATAAATAAATGACCTCTCTTGTCTATTTGGTTTTCGAAAAAG 4055
Db 61 TAGCTTTTAAATAAACAATAAATAAATGACCTCTCTTGTCTATTTGGTTTTCGAAAAAG 120
Qy 4056 ACCCACTATCAAGGATGCTGATGCTGCTATTAAAAATTTGTTCCAAATGTTCCAT 4109
Db 121 ACCCACTATCAAGGATGCTGATGCTGCTATTAAAAATTTGTTCCAAATGTTCCAT 174

RESULT 10
US-08-630-915A-193
; Sequence 193, Application US/08630915A
; Patent No. 6309820
; GENERAL INFORMATION:
; APPLICANT: SPARKS, Andrew B.
; APPLICANT: HOFFMAN, No. 6309820h
; APPLICANT: KAY, Brian K.
; APPLICANT: FOWLES, Dana M.
; APPLICANT: MCCONNELL, Stephen J.
; TITLE OF INVENTION: POLYPEPTIDES HAVING A FUNCTIONAL
; TITLE OF INVENTION: DOMAIN OF INTEREST AND METHODS OF IDENTIFYING AND
; TITLE OF INVENTION: USING SAME
; NUMBER OF SEQUENCES: 227
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Pennie & Edmonds LLP
; STREET: 1155 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: USA
; ZIP: 10036-2711
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; OPERATING SYSTEM: IBM PC compatible
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/630,915A
; FILING DATE: 03-APR-1996
; CLASSIFICATION: 536
; ATTORNEY/AGENT INFORMATION:
; NAME: Mibrock, S. Leslie
; REGISTRATION NUMBER: 18,872
; REFERENCE/DOCKET NUMBER: 1101-174
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 790-9090
; TELEFAX: (212) 869-8864/9741
; TELEX: 66141 PENNIE
; INFORMATION FOR SEQ ID NO: 193:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2873 bases
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: unknown
; MOLECULE TYPE: DNA
US-08-630-915A-193

Query Match 3.2%; Score 165.2; DB 3; Length 2873;
Best Local Similarity 60.4%; Pred. No. 6.4e-29;
Matches 343; Conservative 0; Mismatches 193; Indels 32; Gaps 3;

Qy 3208 GAAATGCCAGGTTATTTGCTCATACACCGCCACCGCCCGCCAGCAGCTCCTCTGCC 3267
Db 1119 GAGATTGCTCAGGTAACCTCAGCATATGTTGCTTCTGTTCTGAACAACTTAGCCTTGA 1178
Qy 3268 CTTGCTCAGCTGATTTTATCCGGAAGAAAGAACCCAGCTGATGTTGGAGGAGAGCTG 3327
Db 1179 CCAGGACAGTTAATATTTAAATTTCTAAAGAAATAACAAGTGGTGGTGGCAAGGAGTTA 1238

HOFFMAN, No. 6709821h
KAY, Brian K.
FOWLES, Dana M.
McCONNELL, Stephen J.
TITLE OF INVENTION: POLYPEPTIDES HAVING A FUNCTIONAL
DOMAIN OF INTEREST AND METHODS OF IDENTIFYING AND
USING SAME
NUMBER OF SEQUENCES: 227
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: USA
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/879,957
FILING DATE: 13-Jun-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,915
FILING DATE: 03-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 1101-174
REFERENCE/DOCKET NUMBER: 1101-174
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864/9741
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 39:
SEQUENCE CHARACTERISTICS:
LENGTH: 747 bases
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 39:
US-09-879-957-39
Query Match 3.2%; Score 163.8; DB 3; Length 747;
Best Local Similarity 63.7%; Pred. No. 6.8e-29;
Matches 277; Conservative 0; Mismatches 137; Indels 21; Gaps 1;
QY 3208 GAAATTGCCAGGTATTGCTCATACACCGCCACCGCCCGCCGAGCAGCTCACTCTCGCC 3267
DB 334 GAGATTGCTCAGGTAACTTCAGCATATGTTGCTTCTGTTCTGACAACTTAGCCTTGCA 393
QY 3268 CCTGGTCAGCTGATTTTGATCCGAAAAAGAACCCAGGTGGATGGTGGGAAGGAGAGCTG 3327
DB 394 CCAGGACAGTTAATTAATTAATTAATTAAGAAAAATACAAAGTGGTGGTGGCAAGGAGTTA 453
QY 3328 CAAGCAGCTGGGAAAAAGCCCGCAGATAGGCTGTTCCAGCTAATTAATGTAAGCTTCTA 3387
DB 454 CAGGCCAGAGGAAAAAGACGACAGAAAGGATGGTTCTCCAGTCATGTTAAACTTTTG 513
QY 3388 AGCCTGGGACGAGCAAAATCACTCCCAACAGAGCCACCTAAGTCAACAGCATTTAGCGCA 3447
DB 514 GGTCAAAGCAGTGAAGAGCCA-----CACCTGCTTTTCATCCT 552
QY 3448 GTGTGCCAGGTGATTTGGGATGTACGACTACACCGCCGAGAAATGACGATAGCTGGCCTTC 3507
DB 553 GTATGTGAGGTGATTTGCTATGATGACTATGACAGCAAAATTAATGAAGATGAGCTCAGTTTC 612
QY 3508 AACAGGCCAGATCATCAAGCTCCTCAACAGGAGGACCTGACTGCTGGTGAAGAGGAA 3567
DB 613 TCCAAGGGACAACTCATTAATGTTATGAACAAAGATGATCCTGATGTTGGTGAAGAGAG 672
US-09-879-957-39
Query Match 3.2%; Score 163.8; DB 3; Length 747;
Best Local Similarity 63.7%; Pred. No. 6.8e-29;
Matches 277; Conservative 0; Mismatches 137; Indels 21; Gaps 1;
QY 3208 GAAATTGCCAGGTATTGCTCATACACCGCCACCGCCCGCCGAGCAGCTCACTCTCGCC 3267
DB 334 GAGATTGCTCAGGTAACTTCAGCATATGTTGCTTCTGTTCTGACAACTTAGCCTTGCA 393
QY 3268 CCTGGTCAGCTGATTTTGATCCGAAAAAGAACCCAGGTGGATGGTGGGAAGGAGAGCTG 3327
DB 394 CCAGGACAGTTAATTAATTAATTAATTAAGAAAAATACAAAGTGGTGGTGGCAAGGAGTTA 453
QY 3328 CAAGCAGCTGGGAAAAAGCCCGCAGATAGGCTGTTCCAGCTAATTAATGTAAGCTTCTA 3387
DB 454 CAGGCCAGAGGAAAAAGACGACAGAAAGGATGGTTCTCCAGTCATGTTAAACTTTTG 513
QY 3388 AGCCTGGGACGAGCAAAATCACTCCCAACAGAGCCACCTAAGTCAACAGCATTTAGCGCA 3447
DB 514 GGTCAAAGCAGTGAAGAGCCA-----CACCTGCTTTTCATCCT 552
QY 3448 GTGTGCCAGGTGATTTGGGATGTACGACTACACCGCCGAGAAATGACGATAGCTGGCCTTC 3507
DB 553 GTATGTGAGGTGATTTGCTATGATGACTATGACAGCAAAATTAATGAAGATGAGCTCAGTTTC 612
QY 3508 AACAGGCCAGATCATCAAGCTCCTCAACAGGAGGACCTGACTGCTGGTGAAGAGGAA 3567
DB 613 TCCAAGGGACAACTCATTAATGTTATGAACAAAGATGATCCTGATGTTGGTGAAGAGAG 672
QY 3568 GTCAATGACAGAGTGGGCTCTCCCATCCAATTAATGTAAGCTGACACAGACATGGAC 3627
DB 673 ATCAAGGGGTGATGGTCTCTTCTTCAACTACGTTAAGATGACGACAGACTCAGAT 732
QY 3628 CCAAGCCAGCAATGA 3642
DB 733 CCAAGTCAACAGTGA 747
RESULT 13
US-09-879-957-39
Sequence 39, Application US/09879957
Patent No. 6709821
GENERAL INFORMATION:
APPLICANT: SPARKS, Andrew B.

